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# BIM geometry with Julia Plasm

Functional language for CAD programming

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 $\mathrm{Pre}/\mathrm{Post}\ \mathrm{conds}\ 4\ \to 5$ 

Example 6

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# Part I Basic Concepts

# Chapter 1 Introduction to Julia Programming

Many programming languages are developed for writing computer programs. Let's imagine their set covered, in a mathematical sense, by a few subsets called *programming paradigms* that characterize the *programming style*, i.e., how the computation is performed, how data are transformed or stored, and other aspects of calculation. Some languages follow a single paradigm; other languages are multiparadigm since their programs may be written adopting more than one style. Examples are Smalltalk (object-oriented), Haskell or ML (functional), Prolog (declarative). C++, Java, Python, and Julia are multiparadigm. Julia was designed for scientific programming, which currently leads for performance, simplicity, and expressiveness. Julia solved the *two-language problem* [?], so unifying program prototyping and optimization. This book begins with a chapter introducing the Julia multi-paradigm which blends high description power and simplicity with very fast computing ability. The chapter is not a programming primer, but is oriented to readers already able to code in other languages.

# 1.1 Basic syntax and type system

Julia combines features of productivity languages, such as Python or MAT-LAB, with characteristics of performance-oriented languages, such as C++ or Fortran. Syntactically, Julia is easy and fast to write and debug, and also enjoys a great collection of packages.

### Basic syntax

Julia strongly resembles Python for generic code, and MATLAB for algebraic calculus with matrices and vectors. Only a few syntax notions are actually necessary to allow the reader to understand the scripts in this book. A set

of differences between the languages Julia, Python, and R, full of scriptlets, may be found in https://cheatsheets.quantecon.org, suggested to readers.

The comment *lines* start with a character #. All text following it on the line is considered a comment and skipped by the compiler. *Multiline* comments, called block-comments, start with #= and terminate with a reverse pair =#.

Even if actually multi-paradigm, Julia can be properly considered a functional language, as everything in Julia is an *expression* generating a *value*.

There are several basic types of numbers: e.g., Int64, Float64, Complex {Float64}, Rational{Int64}. String values are created with "<string chars>"; Char literals are written as 'a','b', etc. All arithmetic infix operators are available, as well as bitwise operators, Boolean values and operations, and primitive comparison operators.

#### Variables

Variables are declared while assigning a value to them. Accessing a non-previously declared variable is an error. *Names* of variables start with a *letter* or *underscore* and may have any number of alphanumeric characters, including  $_{-}$  and  $_{-}$ ! It is also possible to use many Unicode characters, like  $_{\pi}$  or  $_{-}$  using their LATEX names followed by the Tab key, like  $_{-}$   $_{-}$   $_{-}$   $_{-}$ 

Values have a type, but variables do not so that a variable can be reassigned with values of the same or different types.

# Naming conventions

- 1. In long names word separation can be indicated by underscores, but their use is discouraged unless the name would be hard to read otherwise;
- 2. names of Types begin with a capital letter, and word separation is shown with *UpperCamelCase* instead of underscores: AbstractFloat;
- 3. names of functions and macros are in lower case, without underscores;
- 4. functions that modify their input have names that end in !. These functions are called *mutating* functions or *in-place* functions because they mutate their input. For large data structures, this one is rarely a good idea.

### Control flow

Julia provides a variety of control flow constructs:

- 1. Compound Expressions: block begin...end, and ; for expressions in sequence.
- 2. Conditional Evaluation: if...elseif...else...end, and <>?<>:<> (ternary operator).

- 3. Short-Circuit Evaluation: logical operators && ("and"), and  $\parallel$  ("or"), are chained comparisons.
- 4. Repeated Evaluation: Loops: while...end, and for...end.
- 5. Exception Handling: try...catch, error, and throw. Exceptions are thrown when an unexpected condition has occurred.
- 6. Tasks (aka *Coroutines*), a generalized subroutine in Julia, which is a set of instructions stored in memory and invoked during execution. yieldto allow suspending and resuming of tasks.

The for...end loops iterate over iterables; iterable types include Range, Array, Set, Dict, and AbstractString. In loops you can use in instead of range generators

while...end loop lops while a condition *predicate* is true, and terminates when the predicate become false.

Sequential *iteration* is implemented by the iterate function [?]. Instead of mutating objects as they are iterated over, Julia iterators may keep track of the iteration state externally from the object. The return value from iterate is always either a tuple of a value and a state, or **nothing** if no elements remain.

## Type system

Julia has a type system. Every *value* has a type; variables do not have types themselves. You can use the typeof function to get the type of a value: abcd = "abcd"; typeof(abcd) # => String.

Types are first-class values that can be used as arguments of functions and returned by functions. DataType is the type that represents types, including itself.

Julia's type system is *dynamic*. Still, it gains some of the advantages of static type systems by making it possible to indicate that certain values are of specific types and by type reasoning. This can greatly assist in generating efficient code, but even more significantly, it allows *method dispatch* on the types of function arguments to be deeply integrated with the language.

In Julia, a type is *concrete* if it can be instantiated, that is, some type T is concrete if there exists at least one value v such that  $\mathsf{typeof}(v) \# => \top$ . Abstract types cannot be instantiated; they are used to create hierarchies of types, useful to generate fast code. Both abstract and concrete types can be parameterized by other types. They can also be parameterized by symbols and values of types s.t. isbits returns true.

The hierarchical relationships between types are explicitly declared. One particularly distinctive feature of Julia's type system is that *concrete types* may not *subtype each other*: all concrete types are *final*: may only have abstract types as supertypes.

Users can define new types. User-defined types are like *records* or *structs* in other languages. New types are defined using the **struct** keyword. Julia

gives a default constructor as a function with the same name of the type. The default constructor's arguments are the *properties* of the type, in the order they are listed in the definition [?].

These struct-style types are *concrete types*. So, they can be instantiated, but cannot have *subtypes*. The other kind of types is **abstract** types defined as: **abstract** < name > **end**. Abstract types cannot be instantiated, but can have subtypes. AbstractString, as the name implies, is also an abstract type.

#### Type relations and hierarchy

The method supertype(T::DataType) returns the DataType father of T. The *subtype* operator is a predicate T1 <: T2 that returns true when values of type T1 are also of type T2. The *supertype* operator T1 >: T2 is equivalent to T2 <: T1. Two examples follows (see also Figure ??):

```
subtypes # => (generic function with two methods)
subtypes(Number) # => [Complex, Real]|
```

Types are used for documentation, optimizations, and dispatch. In Julia, they are not statically checked, but used by the JIT compiler to create faster code. The :: operator may be used to attach type annotations to expressions and variables within the program. As a general rule, it is better not to use type annotation but let the compiler do it, so writing generic code makes also easier to interact with other packages.

The abstract type declares a type that *cannot* be instantiated and serves only as a node in the type graph, thereby describing sets of *related concrete types*: those concrete types which are their descendants. Abstract types form the conceptual hierarchy, which makes Julia's type system more than just a collection of object implementations. For example: Number has supertype Any, whereas Real is an abstract subtype of Number (see Figure ??).

A primitive type declares a concrete type whose data consists only of a series of bits. Classic examples of primitive types are integers and floating-point values. Some examples of built-in primitive type declarations from Julia implementation:

```
primitive type Char 32 end
primitive type Bool <: Integer 8 end</pre>
```

Type declarations can be used in **global** scope, i.e., type annotations can be added to global variables to make accessing them type stable (about the importance of **type-stability** read [?]).

```
julia> x::Int = 10
10
julia> x = 3.5
ERROR: InexactError: Int64(3.5)
```

#### 1.2 Functions and collections

The actual backbone of Julia programming are the data objects of *collection* types, including arrays, tuples, dictionaries, and sets. Objects of Function type have great significance in the language, since they denote any computable transformation from input data to output results of a computation.

### 1.2.1 Julia functions

In Julia, a function is an object that maps a tuple of argument values to a **return** value. All arguments to functions are passed by sharing (i.e., by memory address). Julia functions are not *pure* mathematical functions because they can alter and be affected by the global state of the program.

### Definition 1.1 (Basic syntax for defining functions)

A generic function can be used for different types of its arguments. In simple words, whenever the function is called with arguments of a new type, the Julia compiler will generate a different version, called *method* for that function. The same happens when a function name is invoked with different number and types of arguments. See Julia Multiple Dispatch [?].

The keyword **function** creates new functions. Functions return the value of their last expression.

**Definition 1.2 (Statement functions)** There is a compact definition of functions, like in old Fortran. Here is the function definition with formal arguments, andt its application to actual argument values:

```
julia> f_add(x,y) = x + y
f_add (generic function with 1 method)
julia> f_add(2,3)
```

**Definition 1.3 (Multiple return values)** Functions can also return multiple values, as a tuple.

```
julia> f(x,y) = x+y, x-y, x*y
f (generic function with 1 method)
julia> a,b,c = f(2,4)
(6, -2, 8)
```

**Definition 1.4 (Variable number of arguments)** You may define functions whose < head > takes a *variable* number of *positional* arguments:

```
function varargs(args...) < body > end
```

of course followed by a body of instructions and by the end token.

The token ... is called a *splat*. We just used it to define a function *head*. The splat can also be used in a *function call*, where it will splat an array or tuple contents into the *argument* list:

```
add(list...) is equivalent to add(5,6,7,8,9) when list = [5,6,7,8,9].
```

**Definition 1.5 (Optional positional arguments)** You may define functions with *optional* positional arguments. with an assigned *default* value, and hence not necessarily with an assigned value at runtime.

The \$\\$ is used to interpolate values of variables or expressions into strings.

**Definition 1.6 (keyword-optional arguments)** You may also define functions that take keyword-optional arguments.

Note the ; character before the optional arguments

Coding 1.2.1 Function call A corresponding function call follows, showing the parameter names k1 and name2 together with the assigned values, used to increase the code readability:

```
julia> keyword_args(; k1=4, name2="hello")
Dict{String, Any} with 2 entries:
    "name2" => "hello"
    "k1" => 4
```

You can combine all kinds of arguments in the same function, with keyword ones defined in the last positions, following the ; character. They can be invoked in any number and order, and possibly substituted by the default value.  $\Box$ 

# Functional programming style

Julia allows the programmer to use efficiently several traits of functional programming style, as it is shown in the following of this section.

*Coding 1.2.2* In other words, Julia has *first class* functions. In the following scriptlet, the function create\_adder returns an adder function:

```
function create_adder(x)
    adder = function(y)
        return x + y
    end
    return adder
end
```

It is also possible to name the internal function if desired:

To be called, e.g., add\_10 = create\_adder(10); add\_10(3) #=> 13.

**Definition 1.7 (lambda syntax)** The *lambda syntax* or "stabby lambda syntax" is used to create *anonymous* functions:  $(x \rightarrow x + 2)(3) \# => 5$ , where the lambda expression is  $x \rightarrow x + 2$ . The *arguments* are before the characters " -> ", and after the stab we have the *value-generating expression*.

```
function create_adder(x)
    y -> x + y
end
```

Of course, it is possible to use a tuple of arguments in the lambda form of functions:  $((x,y) \rightarrow x + y)(2, 3) \# => 5$ . This function is identical to the create\_adder implementation above.

Coding 1.2.3 Even more, like a proper functional language, with curried parameters, we may have:

```
add = (x \rightarrow y \rightarrow x + y);
add(2)(3) # => 5.
```

A curried function is a function which takes multiple parameters one at a time, by taking the first argument, and returning a series of functions which each take the next argument until all the parameters have been fixed, and the function application can complete, at which point, the resulting value is returned. Note that the number of arrows equals the number of applications.

# Julia higher-order functions

The map(), filter(), and reduce() functions are three fundamental higher-order functions that are found in almost every programming language today. In Julia we have this syntax for built-in higher-order functions:

```
map(add_10, [1,2,3]) # => [11, 12, 13]
filter(x -> x > 5, [3, 4, 5, 6, 7]) # => [6, 7]
reduce(*, [2; 3; 4]; init=-1) # => -24
```

Reduce takes two arguments — a function f and a collection A. The function f must take two arguments, and then reduce goes through the collection and one-element-at-a-time it updates result = f(result, elt). The keyword argument init is the initial value to use in the reductions. For +, \*, max, and min the init argument is optional.

```
julia> a = reshape(Vector(1:16), (4,4))
  4×4 Matrix{Int64}:
  1 5 9 13
   2 6 10 14
   3 7
        11
            15
   4 8 12
            16
julia> reduce(max, a, dims=2)
  4×1 Matrix{Int64}:
  13
   14
  15
julia> reduce(max, a, dims=1)
 1×4 Matrix{Int64}:
  4 8 12 16
```

You may also consider using foldl or foldr, with fixed associativity direction: foldr(op, itr; [init]) is like reduce, but with guaranteed right associativity. Or foldl(op, itr; [init]) is like reduce, but with guaranteed left associativity, foldr(\*, 1:5; init=1) # => 120 in this case also foldr(\*, 1:5; init=1) # => 120. Such recursive functionals transform any binary operator into an n-ary operator and might implement many geometric programming patterns.

Remark 1.1 When redefining a method or adding new methods, realizing that these changes don't take effect *immediately* is essential.

This is key to Julia's ability to statically infer and compile code to run fast without the usual JIT tricks and overhead. Indeed, any new method definition won't be immediately visible to the current runtime environment, including Tasks and Threads (and any previously defined <code>@generated</code> functions, an excellent <code>metaprogramming</code> tool [?]).

### 1.2.2 Collections

Julia Base package contains a variety of functions and macros suitable for performing scientific and numerical computing but is as broad as many general-purpose programming languages. Additional functionality is available from a growing collection of available packages.

The proper Julia collections are arrays, tuples, dictionaries, and sets. All (but sets) can be accessed by integer indices. User-defined collections must satisfy the Iterable Collections protocol.

Beware, Julia indexes everything from 1 (like MATLAB and Fortran), not 0 (like most languages, including C, C++, Rust, and Java). Or else, iterating over strings is recommended (map, for loops, etc). The notation \$(expr) can be used for interpolation of a value inside a string, making complex printing very easy. Note that parenthesis () are mandatory when expr is not a single token.

# Arrays

Julia provides a complete collection of basic arithmetic and bitwise operators across all of its numeric primitive types and portable, efficient implementations of a comprehensive collection of standard mathematical functions.

 $\label{like types} AbstractArray\{T,\ N\} \ is the supertype for N-dimensional arrays (or array-like types) with elements of type T; array and other types are subtypes of this.$ 

 $AbstractVector\{T\} \ is the supertype \ for \ one-dimensional \ arrays \ (or \ array-like \ types) \ with \ elements \ of \ type \ T. \ It \ is \ an \ alias \ for \ AbstractArray\{T,1\}.$ 

AbstractMatrix{T} is the supertype for two-dimensional arrays (or array-like types) with elements of type T. It is an alias for AbstractArray{T,2}.

A broadcast dot operator (.) applies any Julia operator or function to all the elements of a collection: f.[a,b,c] # => [f(a),f(b),f(c)].

#### **Tuples**

In Julia, typeof(Tuple) # => DataType is an *immutable* collection of distinct values of the same or different datatypes separated by commas.

Tuples are a *heterogeneous* collection of values. Tuples are more like arrays in Julia, except that arrays only take values from the same datatype. The values in a tuple *cannot be changed* because tuples are immutable. The whole tuple value in a variable can only be *replaced* by a new tuple value.

The sequence of values stored in a tuple can be of any type, and integers index them. Although not needed, defining a tuple using *parentheses* around the sequence of values is expected. This helps in understanding the Julia tuples more easily.

Coding 1.2.4 The tuple function returns a tuple from given objects:

```
tuple(1, 'b', pi) # => (1, 'b', π).
```

**Coding 1.2.5** The function ntuple(f::Function, n::Integer) creates a tuple of length n, computing each element as f(i), where i is the index of the element:

```
ntuple(i -> 2i, 4) # => (2, 4, 6, 8).
```

#### Dictionary

A simple look-up table is a helpful way of organizing many data types: given a single piece of information, such as a number, string, or symbol, called the *key*, what is the corresponding data *value*? For this purpose, Julia provides the Dictionary object, called Dict for short. It is an "associative collection" because it associates keys with values [?].

**Definition 1.8 (Dict is the standard dictionary type)** Its implementation uses hash as the hashing function for the key and **isequal** to determine equality. Redefine these two functions for *custom types* to override how they are stored in a hash table. Any hash function must compute an integer hash code such that isequal(x,y) implies hash(x) == hash(y).

Dictionaries can be created by passing *pair objects* constructed with => to a Dict *constructor*: Dict("A"=>1, "B"=>2). This call will attempt to

infer type information from the keys and values (i.e., this example creates a Dict{String, Int64}. To explicitly specify types use the syntax Dict{KeyType,ValueType}(...).

Coding 1.2.6 Construction syntax directly using pairs key => value:

```
julia> Dict{String, Int32}("A"=>1, "B"=>2) #=
Dict{String, Int32} with 2 entries:
   "B" => 2
   "A" => 1 =#
```

#### Coding 1.2.7 Dictionaries may also be created using generators.

```
julia> f = i->2i; Dict(i => f(i) for i=1:10) #=
Dict{Int64, Int64} with 10 entries:
   5 => 10
   4 => 8
   : => : =#
```

Remark 1.2 Let us notice that the pairs in a Dict are not ordered according to the generation sequence. OrderedDict in DataStructures package is used for this purpose.

Given a dictionary D, the syntax D[x] returns (in reading) the value associated to key x (if it exists) or throws an error, and D[x] = y stores (in writing) the key-value pair  $x \Rightarrow y$  in D (replacing any existing value for the key x).

Multiple arguments to D[...] are converted to tuples; for example, the syntax D[x,y] is equivalent to D[(x,y)], i.e., it refers to the value keyed by the tuple (x,y).

# Set

**Definition 1.9** Sets are *mutable containers* that provide fast membership testing.  $Set{T} <: AbstractSet{T}$ .

Set datatype enjoys efficient implementations of set operations such as in, union, and intersect. Elements in a Set are *unique*, as determined by the elements' definition of isequal. The order of elements in a Set is an *implementation detail* and cannot be relied on.

#### Coding 1.2.8 Some Set examples follow:

```
julia> s = Set("aaBca") #=
Set{Char} with 3 elements:
```

```
1 C 1
 'B'
            =#
julia> push!(s, 'b')
                              #=
Set{Char} with 4 elements:
  'a'
  1 C 1
  'b'
  'B'
            =#
julia> union([4 2 3 4 4], 1:3, 3.0)
4-element Vector{Float64}:
 4.0
 2.0
 3.0
  1.0
```

Let note the type promotion of all Set elements to the same type Float64.  $\square$ 

# 1.3 Matrix computations

A primitive type is a concrete type whose data consists of plain old bits. Classic examples of primitive types are integers and floating-point values. Julia's primitive numeric types called bits types (integers, both signed and unsigned, Booleans, and floats) are Int8, UInt8, Int16, UInt16, Int32, UInt32, Int64, UInt64, Int128, UInt128, Bool, Float16, Float32, Float64, depending on the number of bits. Additionally, full support for Complex and Rational numbers is built on top of these primitive numeric types.

# Vector and Matrix Types

In Julia, the Array type holds both "bits" values as well as "boxed" values. Boxed variables are *heap-allocated* and tracked by the garbage collector.

The distinction is whether the value itself is stored *inline* (in the directly allocated memory of the array, i.e., without following pointers to it) or if the array's memory is simply a collection of pointers to objects allocated elsewhere. Regarding performance, accessing values inline is a substantial advantage over following a *pointer* to the actual values.

Julia provides a first-class array implementation without treating arrays in any special way. The array library is implemented almost entirely in Julia itself and derives its performance from the compiler, just like any other code written in Julia.

The  $type\ Vector\{T\}\ <:\ AbstractVector\{T\}\$ is a 1-dimensional  $dense\ ar-ray$  with elements of type T, often used to represent a mathematical vector. It is an alias for  $Array\{T,1\}$ .

Of course, the method Vector{T}(undef,n) constructs an uninitialized object Array{T,1} of length n. Analogously, we have the Base.Matrix as the alias Matrix in Base package:

- the parameterized type Matrix{T} <: AbstractMatrix{T}, is a two-dimensional dense array with elements of type T, often used to represent a mathematical matrix, where stands for Array{T,2};</li>
- 2. the  $method \, Matrix\{T\} \, (undef, m, n)$  which construct an  $uninitialized \, Array \, \{T, 2\}$  of size  $m \times n$ .

#### Vectors in Julia

We see how to create and manipulate vectors in Julia and how Julia notation differs from mathematical notation.

To create the 3-vector 
$$x = (8, -4, 3.5) = \begin{pmatrix} 8 \\ -4 \\ 3.5 \end{pmatrix}$$
, use:  $x = [8, -4, 3.5]$ ,

but x = [8;-4;3.5] also works. Be careful for similar-looking expressions, because (8,-4,3.5), [8,-4,3.5] and [8-43.5] are not equivalent in Julia. They are a tuple, a column matrix, and a row matrix, respectively.

To get an integer range from i to j (for  $i \leq j$ ), let's use a colon iterator i:j. The assignment x = collect(1:10) returns the array x. To specify an increment size, add an argument.

```
x = collect(1:0.1:10)' #= >
[1.0 1.1 1.2 1.3 1.4 ... 9.4 9.5 9.6 9.7 9.8 9.9 10.0]. =#
```

The range from 1 to 10 with 0.1 step size is given above. Note the apex for transposition, which is needed for typographical reasons here.

- 1. indexes run from 1 to n = length(x), and  $x_2$  is x[2]
- 2. you can also set an element, e.g., x[3] = 10.5;
- 3. use a range to select more than one element; x[2:3] selects the second and third elements;
- 4. x[end] selects the last element;
- 5. to select the even significant elements of x use

```
x[1:2:end]' # => [1.0 1.2 1.4 ... 9.6 9.8 10.0].
```

To form a stacked Vector made by vectors a=[1;2] and b=[3;4;5] use [a...,b...] or [a;b].

The expression [a,b] would return a  $Vector{Int64}$  value with 2 elements of type  $Vector{Int64}$ . Stacked vectors can be used as list of lists. To access an element in W = [a,b] use W[2][2] # => 4

Many more vector operations are defined in Base package. In Julia, a scalar and a vector can be added using the dot (broadcast) operator. The scalar is added to each entry of the vector: [2,4,8] .+ 3 # = [5,7,11].

```
Scalar-vector multiplication uses \ast because the operator is linear:
```

```
[2,4,8] * 3 # => [6,12,24]. Both expressions are commutative.
```

In Julia syntax, like in MATLAB and differently from Python: + and \* operate on congruent arrays, otherwise promote the arguments to same type.

### Matrices in Julia

Julia provides a very simple notation to create matrices. A matrix can be created using the following notation:  $A = [1\ 2\ 3;\ 4\ 5\ 6]$ . Spaces separate entries in a row and semicolons separate rows. We can also get the size of a  $m \times n$  matrix using size(A) = (m,n). Block matrices are easily generated by the same notation, as long as matrix or vector blocks of consistent size are used.

In addition to (and as part of) its support for multi-dimensional arrays, Julia provides native implementations of many common and useful linear algebra operations which can be loaded with using LinearAlgebra. Basic operations, such as tr, det, and inv are all supported in LinearAlgebra package. As well as other useful operations [?], such as finding eigenvalues or eigenvectors:

```
A = [-4. -17.; 2. 2.] #=
2×2 Matrix{Float64}:
-4.0 -17.0
2.0 2.0 =#
eigvals(A) #=
2-element Vector{ComplexF64}:
-1.0 - 5.0im
-1.0 + 5.0im =#
eigvecs(A) #=
2×2 Matrix{ComplexF64}:
0.945905-0.0im 0.945905+0.0im
-0.166924+0.278207im -0.166924-0.278207im =#
```

### 1.4 Linear algebra and sparse arrays

In addition to its support for multi-dimensional arrays, Julia provides native implementations of many common and useful linear algebra operations, which can be loaded by **using** the package. A simple example is given. Note the promotion of values in matrix A:

```
using LinearAlgebra
A = [3. 9 8; 5 9 1; 4 8 2] #=
3x3 Matrix{Int64}:
3.0 9.0 8.0
5.0 9.0 1.0
4.0 8.0 2.0 =#
A' #=
3x3 adjoint(::Matrix{Int64}) with eltype Int64:
3.0 5.0 4.0
9.0 9.0 8.0
8.0 1.0 2.0 =#
```

Basic operations, such as tr, det, rank, and inv are supported by the package LinearAlgebra:

The inner product  $a^{\top}b$  is written as LinearAlgebra.dot(a,b). Vectors a and b must have the same length.

```
The norm x=\sqrt{(x_1^2+x_2^2+\cdots+x_n^2)} is written LinearAlgebra.norm(x). The distance of two vectors dist(x,y)={\bf y}- x is LinearAlgebra.norm(y-x). Of course, the qualification LinearAlgebra. is not explicitly needed, after declaration, for exported package symbols. The function Root Mean Square: rms(x)=\sqrt{((x_1^2+\cdots+x_n^2)/n)}=x/\sqrt{n} can be expressed as:
```

From Rosetta Code [?] we report several ways of implementing/using the rms(x) function. In fact, there are a variety of ways to do this via built-in functions in Julia, given an array A = rand(10) of values.

The formula can be implemented directly using the comma broadcast:

```
sqrt(sum(A.^2)/length(A))
```

or shorter by using Statistics package:  $sqrt(mean(A.^2))$ . The implicit allocation of a new array by  $(A.^2)$  can be avoided by using sum as a higher-

order function:  $sqrt(sum(x \rightarrow x*x, A)/length(A))$ . Of course, one can also use an explicit loop for near—C performance:

```
function rms(A)
  s = 0.0
  for a in A
     s += a*a
  end
  return sqrt(s/length(A))
end
```

Potentially even better is to use the built-in norm function, which computes the square root of the sum of the squares of the entries of A in a way that avoids the possibility of spurious floating-point overflow (if the entries of A are so large that they may overflow if squared): norm(A)/sqrt(length(A)).

To solve a linear system, Ax = b by x = LinearAlgebra.inv(A) \* b is not recommended for big A matrices. A better approach would be to compute  $x = A \setminus b$ , which is equivalent to

```
qr(A, Val(true)) \ b
```

This expression uses a pivoted QR factorization to solve the least-squares problem, which is much more accurate than the normal-equations approach for poorly conditioned A.

In addition, Julia provides many factorizations which can be used to speed up problems such as linear solve or matrix exponentiation by pre-factorizing a matrix into a form more amenable (for performance or memory reasons) to the problem. In factorize(A) documentation [?] we find:

Compute a suitable factorization of A, based upon the type of the input matrix. factorize checks A to see if it is symmetric/triangular/etc. if A is passed as a generic matrix. factorize checks every element of A to verify/rule out each property. It will short-circuit as soon as it can rule out symmetry/triangular structure. The return value can be reused for efficient solving of multiple systems. For example:

```
A = factorize(A); x = A \setminus b.
```

Furthermore,

```
Y = A \ C.
```

Technically, two matrices cannot be divided, but if you remember the basic concept of division of two fractions, then second fraction remains the same but the first fraction gets inverted and multiplied to the former fraction for performing the division. Example with rational numbers:

```
(4//3 \setminus 5//2 == 3//4 * 5//2 == 15//8) == true.
```

#### Sparse Arrays

In numerical analysis and scientific computing, a sparse matrix is a matrix with many zero elements. In other words, the sparse array is an array where many elements have a recurring value, typically zero. It is, therefore, possible to use a better storage model, storing only the non-zero values. There are several storage schemes for numeric sparse vectors and sparse matrices.

The most straightforward scheme is called *storage by triples* (i,j,v), where some suitable data structure supplies the row index, the column index, and the corresponding value of every non-zero. In Julia, this is implemented in the SparseArray package as input method sparse() with input parameters (I, J, V): two arrays of integer indices and one array of bits type for values.

Typical implementation schemes for sparse matrices are the CSR (compressed sparse row) and CSC (compressed sparse column). Julia uses, by now, only the CSC scheme. Julia supports sparse vectors and sparse matrices in the SparseArrays stdlib module. Sparse arrays contain enough zeros that storing them in a special data structure leads to savings in space and execution time compared to dense arrays.

As said above, n Julia, sparse matrices are stored in the *Compressed Sparse Column* (CSC) format. Julia sparse matrices have the type SparseMatrixCSC {Tv, Ti}, where Tv is the type of the stored values, and Ti is the integer type for storing column pointers and row indices.

The internal representation of type SparseMatrixCSC is as follows:

In some applications, storing explicit zeros in a SparseMatrixCSC is convenient. The nnz function returns the number of elements explicitly stored in the sparse data structure, including non-structural zeros. To count the exact number of numerical nonzeros, use zcount(!iszero, x), which inspects every stored element of a sparse matrix. dropzeros, and the in-place dropzeros!, can be used to remove stored zeros from the sparse matrix.

Sparse vectors are stored in a close analog to a compressed sparse column format for sparse matrices. In Julia, sparse vectors have the type SparseVector{Tv, Ti where Tv is the type of the stored values and Ti is the integer type for the indices. The internal representation is as follows:

```
struct SparseVector{Tv,Ti<:Integer} <: AbstractSparseVector{Tv,
    Ti}
    n::Int  # Length of the sparse vector</pre>
```

```
nzind::Vector{Ti}  # Indices of stored values
nzval::Vector{Tv}  # Stored values, typically nonzeros
end
```

The sparse function is often handy for constructing sparse arrays. For example, to construct a sparse matrix, we can input a vector I of row indices, a vector J of column indices, and a vector V of stored values (this is also known as the COO (coordinates) format). sparse(I,J,V) then returns a sparse matrix S such that S[I[k], J[k]] = V[k]. The equivalent sparse vector constructor is sparsevec, which takes the (row) index vector I and the vector V with the stored values and constructs a sparse vector R such that R[I[k]] = V[k].

The inverse of the sparse and sparsevec functions is findnz, which retrieves the inputs used to create the sparse array. findall(!iszero, x) returns the cartesian indices of non-zero entries in x (including stored entries equal to zero).

Details and examples can be found in the Sparse Vectors and Matrices section of the standard library reference [?].

# 1.5 Parallel and distributed computing

Julia makes easier the implementation of complex algorithms by providing native support for concurrency and fast computation by design. Implementing concurrency and multithreading of computations in Julia is mainly a matter of following some style rules and inserting a few macros in the program code.

Parallelism at all levels, from instruction execution to distributed computation, is mostly implicit. Task-based control flows for parallel execution are natively provided, like booth cooperative multitasking and thread-based preemptive multitasking.

Of course, Julia's multithreading model offers the ability to schedule tasks simultaneously on more than one thread or CPU core, sharing memory.

# 1.5.1 Parallel Programming

There are three steps to program parallelization. First, you identify a decomposition into tasks to be computed concurrently.

If the number of tasks is too large compared to the number of threads, the overhead of scheduling them could slow down the computation. No free lunches. The third step is to figure out how to map tasks to threads [?].

According to Julia manual, the language supports four major classes of concurrent and parallel programming [?], also known as models of parallel

computing: (a) asynchronous "tasks", or coroutines; (b) multi-threading; (c) distributed computing; (d) GPU computing.

# Asynchronous tasks (coroutines)

Coroutines are program components that allow execution to be suspended and resumed, generalizing subroutines (computation with effects as opposed to functions returning a value) for *cooperative* multitasking. This one is a style of multitasking in which the operating system never executes a context switch at system level from a running process to another process. Conversely, tasks voluntarily yield control periodically or when idle or logically blocked.

Tasks are a control flow feature that allows computations to be suspended and resumed in a flexible manner. Julia tasks allow suspending and resuming computations for I/O, event handling, producer-consumer processes, and similar patterns, in order to run multiple applications concurrently.

Objects of type Task can be created by the macro <code>@task x</code>, where x is any expression, usually <code>begin; ...; end</code>. The Task object can be assigned to a variable: <code>t = @task x</code>. After creation, the task must be started; it is started by calling <code>schedule(t)</code>, where it is added to a queue of tasks waiting resources for execution. It is common to create a task and <code>schedule</code> it immediately. The macro <code>@async</code> is provided for that purpose: <code>@async x</code> is equivalent to <code>schedule(@task x)</code>.

The macro <code>@async</code> wrap an expression in a <code>Task</code> and add it to the local machine's scheduler queue. Values can be interpolated into <code>@async</code> via <code>\$</code>, which copies the <code>value</code> directly into the constructed underlying <code>closure</code>. This allows you to insert the value of a variable, <code>isolating</code> the asynchronous code from <code>changes</code> to the variable's value in the current task.

It is strongly encouraged to favor Threads.@spawn over @async always. This is because a use of @async disables the migration of the parent task across worker threads in the current implementation of Julia. Thus, seemingly innocent use of @async in a library function can have a large impact on the performance of very different parts of user applications [?].

The macro @sync waits until all lexically-enclosed uses of @async, @spawn, @spawnat and @distributed are complete. In practice it works like a parallel barrier in other languages. One of the simplest examples is using @sync/@async for non-blocking I/O. For example, you want to download 10 web pages. If you do it in a simple blocking loop, most of the time Julia does nothing but waits for network:

```
urls = ["https://discourse.julialang.org/" for i=1:10]
results = Vector(10)
@time for (i, url) in enumerate(urls)
    results[i] = Requests.get(url)
end
```

But network I/O in Julia is non-blocking, so you can use task machinery to accelerate:

```
@sync for (i, url) in enumerate(urls)
    @async results[i] = Requests.get(url)
end
```

In the following example each println request is run in a separate task using Threads.@spawn and then wait for the result of all of them because of @sync. When one of these tasks encounters I/O operation, it gives away control so that other tasks could use CPU. When I/O operations are finished, the task is resumed via @sync.

```
julia> Threads.nthreads() # => 4
```

```
julia> @sync begin
    Threads.@spawn
        println("Thread-id $(Threads.threadid()), task 1")
    Threads.@spawn
        println("Thread-id $(Threads.threadid()), task 2")
end;
# =>
Thread-id 3, task 1
Thread-id 1, task 2
```

Tasks communicate via Channels. While strictly not parallel computing, Julia lets you schedule Tasks on several threads. When a piece of computing work (in practice, executing a particular function) is designated as a Task, it becomes possible to interrupt it by *switching* to another Task. The original Task can later be resumed, at which point it will pick up right where it left off.

At first, this may seem similar to a function call. However there are two key differences. First, switching tasks does not use any space, so any number of task switches can occur without consuming the call stack. Second, switching among tasks can occur in any order, unlike function calls, where the called function must finish executing before control returns to the calling function [?].

Julia tasks are not threads. They are coroutines which can be scheduled asynchronously on a single thread or multiplexed onto a thread pool. All I/O in Julia is non-blocking and yields to the scheduler, so it is more of a cooperative situation.

#### Data parallelism

For data parallelism, a higher-level description is appropriate. It also helps you write more reusable code; e.g., using the same code for single-threaded, multi-threaded, and distributed computing

In particular, it is important to use libraries that help you describe *what* to compute rather than *how* to compute. Practically, it means to use generalized form of map and reduce operations and learn how to express your computation in terms of them. Luckily, if you already know how to write *iterator comprehensions*, there is not much more to learn for accessing a large class of data parallel computations. [?].

Like how multi-threading is setup, you need to setup multiple worker processes to get speedup. You can start julia with -p auto and with -t auto to maximize the number of processes and tasks *sharing memory*.

Mapping is probably the most frequently used function in data parallelism. In sequential code we have: map(f, vect) which results in

```
[f(vect[i] for i in vect]; but Julia's standard library
```

Distributed.jl contains a function pmap as a distributed version of map, so that you can write using Distributed; pmap(f,vect). Also you can use Folds library and write: using Folds; Folds.map(f,vect) with good speedups on large collections.

```
pmap(f, [::AbstractWorkerPool], c...; distributed=true, [...])
```

Transform collection c by applying f to each element using available workers and tasks. For multiple collection arguments, apply f elementwise. Note that f must be made available to all worker processes; see Code Availability and Loading Packages for details. If a worker pool is not specified, all available workers, i.e., the default worker pool is used. By default, pmap distributes the computation over all specified workers. To use only the local process and distribute over tasks, specify distributed=false.

Julia's *iterator comprehension* syntax is a powerful tool for composing mapping, filtering, and flattening. Recall that sequential mapping can be written as an array or iterator comprehension:

```
b1 = map(x -> x + 1, 1:3)  # => [2,3,4]

b2 = [x + 1 \text{ for } x \text{ in } 1:3]  # array comprehension

b3 = collect(x + 1 \text{ for } x \text{ in } 1:3)  # iterator comprehension

@assert b1 == b2 == b3  # => true
```

The *iterator comprehension* can be executed with threads via Folds.collect [?]:

```
b4 = Folds.collect(x + 1 for x in 1:3)
@assert b1 == b4 # => true
```

Functions such as sum, prod, maximum, and all are the examples of *reduction* (aka fold) that can be parallelized. Using Folds.jl, a sum of an iterator created by the comprehension syntax can easily be parallelized by

```
d = Folds.sum(x + 1 for x in 1:3).
```

# Multi-threading

Multithreading enables programmers to speed up their programs by taking advantage of concurrent execution of multiple threads, with each thread assigned to a different CPU core. On the surface, this type of programming may seem easy, but in practice it can be difficult to ensure correctness and to obtain significant speedup.

Julia supports two different models for multithreaded programming: loop parallelism with the <code>@threads</code> macro and task parallelism with the

Threads.@spawn macro, which is low-level basic construct. Julia's multi-threading provides the ability to schedule Tasks simultaneously on more than one thread or CPU core, sharing memory. This is usually the easiest way to get parallelism on one's PC or on a single large multi-core server. Julia's multi-threading is composable. When one multi-threaded function calls another multi-threaded function, Julia will schedule all the threads globally on available resources.

Although Julia's threads can communicate through shared memory, it is notoriously difficult to write correct and data-race free multi-threaded code. Julia's Channels are thread-safe and may be used to communicate safely. The best way to ensure data-race freedom is to acquire a lock around any access to data that can be observed from multiple threads.

By default, Julia starts up with a single thread of execution.

Threads.nthreads() returns the number of thrads, setted by parameter -t when starting Julia, for example -t auto or --threads n. The function Threads.threadid() returns the integer id of the current thread.

Julia supports parallel loops using the Threads. @threads macro. This macro is affixed in front of a for loop to indicate to Julia that the loop is a multi-threaded region. Julia supports accessing and modifying values atomically, that is, in a thread-safe way to avoid race conditions.

When a program's threads are busy with many tasks to run, tasks may experience delays which may negatively affect the responsiveness and interactivity of the program. To address this, you can specify that a task is interactive.

External libraries, such as those called via ccall, pose a problem for Julia's task-based I/O mechanism. If a C library performs a blocking operation, that prevents the Julia scheduler from executing any other tasks until the call returns.

There are a few specific limitations and warnings to be aware of when using threads in Julia [?].

# 1.5.2 Multiprocessing and Distributed Computing

Most modern computers possess more than one CPU, and several computers can be combined together in a cluster. Harnessing the power of these multiple CPUs allows many computations to be completed more quickly.

There are two major factors that influence performance: the speed of the CPUs themselves, and the speed of their access to memory. In a cluster, it's fairly obvious that a given CPU will have fastest access to the RAM within the same computer (node). Perhaps more surprisingly, similar issues are relevant on a typical multicore laptop, due to differences in the speed of main memory and the cache [?].

Distributed computing runs multiple Julia processes with separate memory spaces. These can be on the same computer or multiple computers. The Distributed standard library provides the capability for remote execution of a Julia function. With this basic building block, it is possible to build many different kinds of distributed computing abstractions. Packages like DistributedArrays.j are an example of such an abstraction. On the other hand, packages like MPI.jl and Elemental.jl provide access to the existing MPI ecosystem of libraries.

An implementation of distributed memory parallel computing is provided by module Distributed as part of the standard library shipped with Julia.

Consequently, a good multiprocessing environment should allow control over the "ownership" of a chunk of memory by a particular CPU. Julia provides a multiprocessing environment based on message passing to allow programs to run on multiple processes in separate memory domains at once [?].

Distributed programming in Julia is built on two primitives: remote references and remote calls. A remote reference is an object that can be used from any process to refer to an object stored on a particular process. Remote references come in two flavors: Future and RemoteChannel. A remote call is a request by one process to call a certain function on certain arguments on another (possibly the same) process.

Much easier is parallel programming in Julia by making use of

DistributedArrays.jl. This great computational abstraction uses the stdlib Distributed to implement a Global Array interface. In fact, large computations are often organized around large arrays of data. In these cases, a particularly natural way to obtain parallelism is to distribute arrays among several processes.

This combines the memory resources of multiple machines, allowing use of arrays too large to fit on one machine. Each process can read and write to the part of the array it owns and has read-only access to the parts it doesn't own. This provides a ready answer to the question of how a program should be divided among machines.

A DArray is distributed across a set of workers. Each worker can read and write from its local portion of the array and each worker has read-only access to the portions of the array held by other workers. Julia distributed arrays are implemented by the DArray type. A DArray has an element type and dimensions just like an Array. A DArray can also use arbitrary array-like types to represent the local chunks that store actual data. The data in a DArray is distributed by dividing the index space into some number of blocks in each dimension.

By using DistributedArrays, common kinds of arrays can be constructed with distributed data structures. E.g., d = DistributedArrays; d.zeros, d.ones, d.rand, d.randn, d.fill. The constructor that buils a distributed array is:

```
DArray(init, dims, [procs, dist])
```

- 1. the parameter init is a function that accepts a tuple of index ranges. This function should allocate a local chunk of the distributed array and initialize it for the specified indices.
- 2. dims is the overall size of the distributed array.
- 3. procs optionally specifies a vector of process IDs to use. If unspecified, the array is distributed over all worker processes only. Typically, when running in distributed mode, i.e., nprocs() > 1, this would mean that no chunk of the distributed array exists on the process hosting the interactive julia prompt.
- 4. dist is an integer vector specifying how many chunks the distributed array should be divided into in each dimension.

For example, the dfill function that creates a distributed array and fills it with a value v is implemented as:

```
dfill(v, args...) = DArray(I->fill(v, map(length,I)), args...)
```

# 1.5.3 Programming the GPU

While kernel functions for GPU are usually written in a C/C++ dialect, the Julia GPU compiler provides the ability to run Julia code natively on GPUs. There is also a rich ecosystem of Julia packages that target GPUs. The JuliaGPU.org website provides a list of capabilities, supported GPUs, related packages and documentation. Julia has several packages for programming GPUs, each of which support various programming models.

JuliaGPU is a Github organization created to unify the many packages for programming GPUs in Julia, in order to perform high-performance GPU programming in a high-level language.

JuliaGPU is vendor-neutral, and some content is available for all supported GPU back-ends. Of course JuliaGPU embraces also vendor-specific tools and APIs. At 2021 JuliaCon workshop, they demonstrated the use of three major GPU programming packages: CUDA.jl for Nvidia GPUs, AMDGPU.jl for AMD GPUs, and oneAPI.jl for Intel GPUs. The various approaches for programming GPUs with these packages, range from generic array operations that focus on ease-of-use, to hardware-specific kernels for when performance matters.

GPU functions (*kernels*) are inherently parallel, so writing GPU kernels is at least as difficult as writing low-level parallel CPU code, but the difference in hardware adds quite a bit of complexity. Most algorithms will need arrays to manage all their data, which calls for a good GPU array library.

# GPUArrays.jl

GPUArrays.jl is the counterpart of Julia's AbstractArray interface, but for GPU types. It is not intended for the end-user, which should only use one of the packages that builds on GPUArrays.jl, such as CUDA.jl (Nvidia), oneAPI.jl (Intel), AMDGPU.jl (AMD), or Metal.jl (Apple), for different hardware support.

GPUArrays is an abstract interface for GPU computations. Think of it as the AbstractArray interface in Julia Base but for GPUs. It allows you to write generic julia code for all GPU platforms and implements common algorithms for the GPU. Like Julia Base, this includes BLAS wrapper, FFTs, maps, broadcasts and mapreduces. So when you inherit from GPUArrays and overload the interface correctly, you will get a lot of functionality for free.

It is important to remark that Julia allows you to write both GPU kernels and surrounding code in Julia itself, while running on most GPU hardware.

In complex analysis, the *Julia set* [?] of a holomorphic function consists of all those points whose behavior after repeated iterations of the function is chaotic, in the sense that it can change drastically following a small initial perturbation. As one can see in [?], the computational example of *Julia set* with Julia (!) strongly motivates why one should move big array computations to the GPU. For large arrays one gets a solid 60-80x speed-up by moving the calculation to the GPU. Getting this speed-up was as simple as converting the Julia array to a GPUArray.

#### CUDA and OpenCL

There is a model gap between CUDA and OpenCL parallel programming, which are the dominant frameworks used to write low-level GPU code. OpenCL is a heterogeneous programming platform that allows applications to run across multiple platforms, including CPUs, GPUs, and other specialized hardware. Conversely, CUDA is a software framework specifically designed to run computations on Nvidia's GPUs.

While CUDA.jl only supports Nvidia hardware, OpenCL.jl supports all hardware but "is a bit rough around the edges". One needs to decide what to use, and will get pretty much stuck with that decision [?].

One might think that the GPU performance suffers from being written in a dynamic language like Julia, but Julia's GPU performance should be pretty much on par with the raw performance of CUDA or OpenCL. Tim Besard, the creator of Julia GPU compiler, did a great job at integrating the LLVM Nvidia compilation pipeline to achieve the same or sometimes even better performance as CUDA C code [?].

#### CUDA programming in Julia

The CUDA.jl package is the main entry point for programming NVIDIA GPUs in Julia. The package makes it possible to do so at various abstraction levels, from easy-to-use arrays down to hand-written kernels using low-level CUDA APIs. The following is synthesized from Reference [?] that you are invited to read.

Julia has first-class support for GPU programming: you can use high-level abstractions or obtain fine-grained control, all without ever leaving your favorite programming language. The purpose of this tutorial is to help Julia users take their first step into GPU computing. In this tutorial, you'll compare CPU and GPU implementations of a simple calculation, and learn about a few of the factors that influence the performance you obtain.

We can perform GPU computations at a high level using the CuArray type, without explicitly writing a kernel function:

```
using CUDA
x_d = CUDA.fill(1.0f0, N) # vector stored on GPU filled of 1.0 (
    Float32)
y_d = CUDA.fill(2.0f0, N) # vector stored on GPU filled of 2.0
```

Here the d means device, in contrast with host. Now let's do the increment:

```
y_d .+= x_d
@test all(Array(y_d) .== 3.0f0) # => Test Passed
```

The statement Array(y\_d) moves the data in y\_d back to the host for testing. If we want to benchmark this, let's put it in a function:

```
function add_broadcast!(y, x)
    CUDA.@sync y .+= x
    return
end

add_broadcast! # => (generic function with 1 method)
@btime add_broadcast!($y_d, $x_d) # => 67.047 µs
    (84 allocations: 2.66 KiB)
```

The most interesting part of this is the call to CUDA.@sync. The CPU can assign jobs to the GPU and then doing other stuff (such as assigning more jobs to the GPU) while the GPU completes its tasks. Wrapping the execution in a CUDA.@sync block will make the CPU block until the queued GPU tasks are done, similar to how Base.@sync waits for distributed CPU tasks. Without such synchronization, you'd be measuring the time takes to launch the computation, not the time to perform the computation. But most of the time you don't need to synchronize explicitly: many operations, like copying memory from the GPU to the CPU, implicitly synchronize execution.

For that particular hardware used, the GPU computation was significantly faster than the single-threaded CPU computation, and the use of multiple CPU threads makes the CPU implementation competitive.

The high-level functionality of CUDA often means that you don't need to worry about writing kernels at a low level. However, there are many cases where computations can be optimized using low-level manipulations [?].

# 1.6 Modules and packages

A Julia module is a named sequence of julia code, typically contained in a file with the the same name:

```
module NameOfModule; <some code>; end # => Main.NameOfModule
```

A module start with the reserved word **module** followed by *<NameOfModule>* and is terminated by **end** word. Modules in Julia help organize code into coherent units and have the following features.

Modules are *separate namespaces*, each introducing a new *global scope*. This is useful, because it allows the same name to be used for different functions or global variables without conflict, as long as they are in separate modules.

Modules have facilities for detailed namespace management: each defines a set of names it exports, and can import names from other modules with using and import. Modules can be precompiled for faster loading, and may contain code for runtime initialization. Typically, in larger Julia packages you will see module code organized into program files [?]. One can have multiple files per module, and multiple modules per file. include behaves as if the

contents of the source file were evaluated in the global scope of the including module.

The recommended style is not to indent the body of the module, since that would typically lead to whole files being indented. Also, it is common to use UpperCamelCase for module names (just like types),

A software library is a suite of data and programming code that is used to develop software programs and applications. In Julia a library is called package. A Julia package contains modules, tests, and documentation. It extends core Julia functionality. You can share your code with the community by developing a package. You can create a Julia package using as support the built-in package manager PkgDev.jl or the package PkgTemplates.jl. The second is easier for a novice.

A good tutorial about how a user may develop a package is [?]. The reader is warmly solicited to see, and try the suggested step-by-step development.

The Julia ecosystem contains over 9,000 packages that are registered in the General registry, which means that finding the right package can be a challenge. Fortunately, there are services that can help navigate the ecosystem, including:

- 1. **JuliaHub**: service that includes search of all registered open source package documentation, code search, and navigation by tags/keywords.
- 2. Julia Packages: to browse Julia packages, filter by categories, and sort them by popularity, creation date or date of last update. Also supports browsing package

# Chapter 2 The Package Plasm.jl

Plasm is a software package for advanced geometry programming in Julia. In particular, it is oriented to generate engineering and architecture models whose computer representations may be classified as maps of topological polyhedra. This is a general domain of shapes, including linearized approximations of curved objects and, of course, the cellular complexes of simplicial, cubical, and polyhedral cells. Summing up, Plasm.jl is an open-source porting to Julia of the original PlasM, an extension to the geometry of Fl language at Function Level by Backus and his group at IBM Almaden in the 1990s. Written originally in Common Lisp, it was then ported to Scheme and C++, then to Python, and now to Julia.

# 2.1 Backus' functional programming

John W. Backus directed the IBM team that invented and implemented the FORTRAN (Formula Translation) language. FORTRAN was the first high-level scientific and technical computer language developed by IBM in 1954-1956 for scientific and engineering applications and subsequently came to dominate scientific computing. FORTRAN is used up to now on supercomputers.

# FP (Functional Programming) and FL (Function Level)

The 1977 ACM Turing Award, in honor and recognition of Turing's contribution to the field of computing, was presented to John Backus at the ACM Annual Conference in Seattle. In introducing the recipient, Jean E. Sammet, Chairman of the Awards Committee, said that it was for Fortran and the BNF (Backus Normal Form) that he was receiving that year's Tur-

ing Award. Therefore, everybody was expecting Backus would describe the work done to implement FORTRAN at IBM in the 1950s.

Conversely, the Backus' Turing Lecture was entitled "Can Programming Be Liberated from the von Neumann Style? A Functional Style and Its Algebra of Programs" [?]. This lecture was enormously influential, and opened a decade and half of renewed research work about functional languages.

#### Figure 1: Primitive Operations and Notation.

```
identity
                           id:x = x
                           err:x = \bot
selectors si
                           s3: \langle x_1, ..., x_n \rangle = x_3
empty test
                           isnull:x = true iff
                                x = < >
                           tl: < x_1, ..., x_n >
                                 = \langle x_2, ..., x_n \rangle
append to left
                           al: \langle x, \langle y_1, ..., y_n \rangle \rangle
                                = < x, y_1, ..., y_n >
                             x:y=x \ (y \neq 1 \ in \ FP.)
constant
                           [f_1, ..., f_n]:x
construction
                                 = < f_1:x, ..., f_n:x >
composition
                            (f \circ g): x = f: (g:x)
                                • f; g):x =
                                f:x \text{ if } p:x = true
                                g:x \text{ if } p:x = false
                                 L otherwise
apply to all
                            \alpha: f: \langle x_1, ..., x_n \rangle
                                  = < f:x_1, ..., f:x_n >
catenate
                                 <\!< x_1... >, < ... x_n >>
                                 = < x_1, ..., x_n >
infix cat
                           f ++ g = cat \circ [f, g]
```

Fig. 2.1 The primitive operations (combining forms) and notations [?] of FP programs. The semantics of FP is embodied in an underlying algebra of programs, a set of function level equalities that may be used to transform programs and to reason about them.

A simple functional programming (FP) system is described in [?]. It is based on combining forms for building programs from simpler programs. An algebra of programs is described whose variables denote FP functional programs and whose "operations" are FP functional forms, the combining forms of FP programs, to support the function-level programming paradigm. It allows building programs from a set of generally useful primitives and avoiding named variables. Only programs, aka functions, may be named. In Figure 2.4 we display the algebraic rules of FP.

FL (short for "Function Level") is a programming language created by IBM Research at Almaden. FL is the result of the effort [?] to design a practical functional programming language based on Backus' FP. FL is intended to be a programming language in which it is easy to write clear, concise, and efficient programs. FL is designed around a rich set of functionals, forms for combining existing programs to construct new ones. This emphasis on programming at the function level results in programs that have a rich mathematical structure useful in reasoning about

and optimizing programs [?]. A short description of FL combining forms and primitive functions is given below, where its geometric extension with Julia's functional syntax is discussed.

# 2.2 FL-based PLaSM in Julia syntax

Here we gives a brief informal description of the Plasm language [?, ?], together with few examples. The section is intended to acquaint the reader with the syntax and style of Plasm programs without going into details. The interpreter and interactive GUI for PLASM were developed at the Sapienza University of Rome in the 1990s, extending the FL syntax and semantics with the only addition of a geometric type called HPC (hierarchical polyhedral complex) and supporting the FL programming style for geometric computing within the wide domain of Computer-Aided Design. The book GP4CAD (Geometric programming for CAD) [?] provides some hundreds of small, simple programs using the FL syntax.

In the past decade, the command-line user interface (CLI), the geometric computational kernel, and the interactive visualizer of "classic" Plasm were ported to Python [?] and C++, respectively, mainly using the functional features of such languages [?]. Analogously, in recent times, a native and extended port to Julia is being carried out [?], used in this book, and briefly described in the following, abridged in a few general points.

- a. Function application and binary function composition are native in Julia, which provides f(x) and  $g \circ f$ , respectively.
- b. FL sequence  $\langle x_1, x_2, \dots, x_n \rangle$  is implemented as array  $[x_1, x_2, \dots, x_n]$ .
- c. All primitive functions (programs) are *pure* (without side effects) and written *uppercase*, while named expressions have capitalized names.
- d. Each Plasm function is *unary*, possibly using an array of arguments.
- e. The geometric types Hpc and Lar are extended with an optional Julia's dictionary of properties.

The point d. is sometime relaxed in Julia Plasm, to reduce the visual rumor. From [?], where the interested reader may find many codes discussed in this book in their FL version, we recall that significant advantages are obtained with this approach in the style and efficiency of program development. More generally, it is well known that functional programming enjoys several good properties:

- The set of syntax rules of a functional language is very small.
- Each rule is very simple.
- The program code is terse and clear.
- The meaning of a program is well understood, since there is no state.
- Functions may be used both as programs and as data.

Programs are easily connected by concatenation and nesting.

## Programs are functions

Generally speaking, a Plasm program is a function. When applied to some input argument, a program produces some output value. Two programs are usually connected by using functional composition, so that the output of the first program is used as input to the second program. Starting from here we use the Julia syntax.

# Program composition and application

The composition of Plasm functions, i.e., FL programs with Julia syntax, works exactly as the composition of mathematical functions. E.g., the application of the composite mathematical function  $f \circ g$  to the x argument

$$(f \circ g)(x) \equiv f(g(x))$$

means that the function g is first applied to x and that the function f is then applied to the value g(x). The Julia Plasm notation for the previous expression is:

$$(f \circ g)(x) \equiv f(g(x))$$

where  $\circ$  stands for binary function composition and g(x) stands for application of the function g to the argument x. Both notations are Julia's native.

## Naming objects

In Plasm, a name can be assigned to every value generated by the language, by using an (unmutable) definition construct, either with or without explicit parameters. In both cases the so-called body of the definition, i.e. the expression which follows the definition head, at the right hand of the "=" symbol, will describe the computational process which generates the value produced by the computation. The parameters which it implicitly/explicitly depends on may be embedded in such a definition. For example, we may have:

```
object = (Fun3 ∘ Fun2 ∘ Fun1)(params);
```

The computational process which produces the object value can be thought as the computational pipeline shown in Figure 2.2.

In this example, the reliance of the model on the parameters is implicit. To modify the generated object value it is necessary (a) to change the source code in either the body or the local environment of its generating



Fig. 2.2 Example of computational pipeline

function; (b) to compile the new definition; and (c) to evaluate again the object identifier.

# Parametrized objects

A parametric geometric model can be defined, and easily combined with other such models, by using a generating function with formal parameters. Such kind of function may be instanciated with different actual arguments, thus obtaining different output values. For example, we may have:

```
object(params) = (Fun3 \circ Fun2 \circ Fun1)(params);

obj1 = object([p_1, p_2, ..., p_n];

obj2 = object([q_1, q_2, ..., q_n];
```

It is interesting to note that the generating function of a geometric model may accept parameters of *any* type, including other geometric objects.

# 2.3 Geometric Programming at Function Level

As we already know, Plasm is a geometry-oriented extension of a subset of the FL language [?, ?], which is a pure functional language based on combinatory logic [Wikipedia]. In particular, the FL language makes use of both pre-defined and user-defined *combinators*, i.e. higher-order functions which are applied to functions to produce new functions. The small but very significant FL subset which is used as the base environment of Plasm is summarized in this section.

Notice that here and in the remainder of this book the infix symbol  $\equiv$  is normally used to tell the reader that the *expression* on its left side evaluates to the *value* on its right side. Sometimes this symbol is also used to denote an equivalence between syntactical forms.

# Elements of FL syntax in Julia

Primitive FL *objects* are characters, numbers and truth values. Primitive objects added to it by Plasm are Hpc and Lar geometric values, discussed in following chapters. Primitive objects, functions, applications and se-

quences are *expressions*. Sequences are expressions separated by commas and contained within a pair of square brackets (Julia Vector type):

```
[5, fun]
```

An application expression exp1(exp2) applies the function resulting from the evaluation of exp1 on the argument resulting from the evaluation of exp2. Julia allows some binary functions to be used both in infix and prefix form:

$$1 + 3 \equiv +(1,3) \equiv 4$$

Application associates to left, i.e. a sequence of repeated applications is evaluated from left to right. Note that this is only possible if all the applications, but possibly the last one, generate a new function to be applied to the next argument:

$$f(g)(h) \equiv (f(g))(h)$$

Application binds stronger than composition, i.e. applications are evaluated first before compositions, as it is shown in the following example. Of course the application of f must generate a function value:

$$f(g) \circ h \equiv (f(g) \circ h)$$

# Combining forms and functions

The function level approach to programming of FL emphasizes the definition of new functions by combining existing functions in various ways. The result of this approach is a programming style based on function-valued expressions. Some important FL *combining forms* and functions follow.

# Construction

The combining form CONS allows application of a sequence of functions to an argument, so producing a sequence of applications:

```
CONS([f_1,\ldots,f_n])(x) \equiv [f_1(x),\ldots,f_n(x)]
```

A CONSed sequence of functions is a sort of *vector function*, that can be composed with other functions and that can be applied to data. E.g., CONS([sin,cos,tan,atan]) when applied to the argument  $\pi$  returns the sequence of applications

```
CONS([sin,cos,tan,atan])(π) #=
4-element Vector{Float64}:
0.0
-1.0
0.0
1.2626272556789115 =#
```

# Apply-to-all

The combining form AA has a symmetric effect, i.e. it applies a function to a sequence of arguments giving a sequence of applications. It is equivalent to the functional map of other languages, Julia included:

```
\mathsf{AA}(\mathsf{f})([\mathsf{x}_1,\ldots,\mathsf{x}_n]) \equiv \mathsf{map}(\mathsf{f}, [\mathsf{x}_1,\ldots,\mathsf{x}_n]) \equiv [\mathsf{f}(\mathsf{x}_1),\ldots,\mathsf{f}(\mathsf{x}_n)]
```

For example, we may apply the trigonometric SIN function to all the elements of a list of numeric expressions:

```
AA(SIN)([0, \pi/3, \pi/6, \pi/2])

\equiv [SIN(0), SIN(\pi/3), SIN(\pi/6), SIN(\pi/2)]

\equiv [0, 0.8660254037844382, 0.499999999999956, 1.0];
```

The reader should notice that numeric computations often introduce round-off and approximation errors. Just remember that  $\pi$  is an irrational number and cannot be represented exactly by using finite precision arithmetic. Also, functions like SIN are computed by using some truncated series expansion.

## *Identity*

The function returns its argument unchanged

```
ID(x) \equiv x
```

In other words, the application of the identity function to *any* argument, gives back the same argument:

```
\begin{array}{l} ID(0.5) \equiv 0.5 \\ ID(SIN) \equiv SIN \\ ID(SIN)(0) \equiv SIN(0) \equiv 0 \end{array}
```

# Constant

The combining form K is evaluated as follows, for whatever  $x_1$  and  $x_2$ :

```
K(x_1)(x_2) \equiv x_1
```

In other words, the first application returns a constant function of value  $x_1$ , i.e. such that when applied to any argument  $x_2$ , always returns  $x_1$ . Some concrete examples follow:

```
K(0.5) \equiv Anonymous-Function K(0.5)(10) \equiv 0.5 K(0.5)(100) \equiv 0.5 K(0.5)(SIN) \equiv 0.5
```

### Composition

The binary composition of functions COMP, denoted in Julia by the symbol "o", is defined in the standard mathematical way, as we already know:

```
(f \circ g)(x) \equiv f(g(x))
```

where • is obtained via the LATEX expression \circ followed by TAB character. This important typing mechanism is standard in Julia and allows the program code to use Greek letters and many mathematical symbols. The *n-ary composition* of functions is also allowed:

```
COMP([f, g, h])(x) \equiv (f \circ g \circ h)(x) \equiv f(g(h(x)))
```

In the following we have, where  $\pi$ , COS and ACOS are the Plasm denotations for Julia's  $\pi$ , cos and acos <sup>1</sup>, respectively:

```
\begin{array}{ll} (\text{ACOS} \, \circ \, \text{COS})(\pi) \, \equiv \, \text{ACOS}(\text{COS}(\pi)) \, \equiv \, \text{ACOS}(-1) \, \equiv \\ & 3.141592653589793 \\ (\text{COS} \, \circ \, \text{ACOS})(-1) \, \equiv \, \text{COS}(\text{ACOS}(-1)) \, \equiv \, \text{COS}(3.141592653589793) \, \equiv \\ & -1 \\ \text{COMP}([\text{acos, cos, acos}])(-1) \, \equiv \, \text{ACOS}(\text{COS}(\text{ACOS}(-1))) \, \equiv \\ & 3.141592653589793 \end{array}
```

### Conditional combinator

This combinator has the following semantic: "IF the predicate p applied to object x is true, THEN apply f to x; ELSE apply g to x". This construct is very useful when it is necessary to apply different actions to input data depending on the value of some predicate evaluated on them, and is possibly more "natural" than the conditional statements available in other languages.

<sup>&</sup>lt;sup>1</sup> Which can be directly used in the **PLASM** code, of course.

Formally, the conditional form IF([p, f, g]) is evaluated as follows:

```
IF([p, f, g])(x)
\equiv f(x) \text{ if } p(x) \equiv TRUE
\equiv g(x) \text{ if } p(x) \equiv FALSE
```

From a syntax viewpoint, we remark that the IF operator is a higher-order function that *must* be applied to a *triplet of functions* in order to return a function which is in turn applied to the input data.

A predicate is a function  $p: T \to \{true, false\}$  where T is a Type. Both true and false are called truth values, and in Julia are Bool values. The predicate p is a function, as well f and g, to be alternatively executed depending on the truth value of the logical expression p(x). E.g., we have:

```
IF([ISINTPOS, K(true), K(false)])(1000) \equiv true IF([ISINTPOS, K(true), K(false)])(-1000) \equiv false
```

where ISINTPOS is a predefined predicate that returns true when applied to some positive integer.

Insert Right/Left

The combining forms INSR and INSL allow the user to apply a binary function f, with signature<sup>2</sup> f:  $D \times D \to D$ , on a sequence of arguments of any length n. In other words, implicitly it is: INSR(f):  $D^n \to D$ . Note that in the right-hand expressions below, f is always applied esplicitly to a pair of arguments:

```
INSR(f)([x_1, x_2,..., x_n]) \equiv f([x_1, INSR(f)([x_2,..., x_n])])
INSL(f)([x_1,..., x_{n-1}, x_n]) \equiv f([INSL(f)([x_1,..., x_{n-1}]), x_n])
```

An interesting use example of the INSL combinator is given below, where the function BIGGER:  $Num \times Num \rightarrow Num$  is defined. The BIGGER function returns the maximum of two arguments; the BIGGEST:  $Num^n \rightarrow Num$  does the same from a list of arguments of  $arbitrary\ length$ :

```
BIGGER # predefined function
BIGGEST = INSL(BIGGER)
SMALLER # predefined function
SMALLEST = INSL(SMALLER)

BIGGER([-10, 0]) = 0
BIGGEST([-10, 0, -100, 4, 22, -3, 88, 11]) = 88
```

<sup>&</sup>lt;sup>2</sup> The signature of a function f from a domain A to a codomain B is the ordered pair of sets (A, B). It is normally associated to f by writing  $f: A \to B$ .

Catenate

The CAT function appends to the first one any number of input sequences, so creating a single output sequence:

```
CAT([[10,30,20],[11],[-7,8,12]]) \equiv [10,30,20,11,-7,8,12])
```

A pair of concrete examples of how the CAT function is used follows. The second one is quite interesting: it gives a *filter* function used to select the non-negative elements of a number sequence:

```
CAT([[10,30,20],[11],[-7,8,12]]) == [10,30,20,11,-7,8,12]

#=

true =#

(CAT o AA(IF([ LT(0), K([]), ID ])))([-101,23,-37.02,0.1,84])

= CAT([ [], [23], [], [0.1], [84] ])

= [23, 0.1, 84]
```

It is useful to abstract a filter function with respect to a predicate and to an argument sequence, by showing this function semantics where curried LE stands for less or equal to its first argument. LT, GE, GT are similar.

```
FILTER(predicate)(sequence)
FILTER(LE(0))([-1,0,1,2,3,4]) == [-1, 0] # => true
```

Distribute Right/Left

The functions DISTR and DISTL are defined as:

```
DISTR([[a,b,c], x]) \equiv [[a,x], [b,x], [c,x]]
DISTL([x, [a,b,c]]) \equiv [[x,a], [x,b], [x,c]]
```

They accordingly transform a *pair*, constituted by an arbitrary expression and by an arbitrary sequence, into a *sequence of pairs*.

# Script 2.3.1

Let us give an example of  $\sf Plasm$  use. The Euler number e is defined as the sum of a series of numbers. In particular:

$$e = \frac{1}{0!} + \frac{1}{1!} + \frac{1}{2!} + \dots + \frac{1}{n!} + \dots$$

We compute an *approximation* of e, named euler, as the sum of the first 21 terms of the series. The factorial function is native in Julia:

```
euler = (ADD o AA(DIV) o DISTL)([1, AA(factorial)(0:20)])
```

The number 20 is the highest positive integer for which the expression factorial(20) does not overflow out of memory assigned to an Int64 number. In Julia, you can use a type BigInt with the big function, converting a number to a maximum precision representation. Hence, redefine factorial as the FACT function below, which uses the ternary conditional statement:

Coding 2.3.1 (Computation by substitution) The euler value is computed here by successive substitutions. Of course, the Julia's optimizing compiler might do a much better job:

```
euler = (ADD o AA(DIV) o DISTL)([1,AA(FACT)(0:9)])

\( \text{(ADD o AA(DIV) o DISTL)}([1, AA(FACT)(0:9)]) \)
\( \text{(ADD o AA(DIV) o DISTL)}([1, [FACT(0), FACT(1), ..., FACT(9)])] \)
\( \text{(ADD o AA(DIV) o DISTL)}([1, [1,1,2,6,...,40320,362880]]) \)
\( \text{(ADD o AA(DIV)}(DISTL([1, [1,1,2,6,...,40320,362880]]) \)
\( \text{(ADD o AA(DIV)}([[1,1], [1,1], [1,2], [1,6],..., [1,362880]]) \)
\( \text{(ADD o AA(DIV)}([[1,1], [1,1], [1,2], [1,6],..., [1,362880]]) \)
\( \text{(ADD o ADD([ 1/1, 1/1, 1/2, 1/6, ..., 1/40320, 1/362880]) } \)
\( \text{(ADD o ADD([ 1/1, 1/1, 1/2, 1/6, ..., 1/40320, 1/362880]) } \)
```

Above, we have seen our first substantial example of Plasm, aka FL computation, as a sequence of expression transformation using the rules of combinators. The round brackets induce the order of transformations included into an expression and often corresponds to applications. The sub-expressions nested more deeply are transformed first. When using Float64, i.e., 8 bytes, the numeric precision is 15-16 digits.

A simpler and more elegant implementation of the Euler number is given below, where C is the currying combinator<sup>3</sup>:

```
EULER(n) = (ADD \circ AA(C(DIV)(1) \circ FACT))(0:n)
```

The best Julia approximation of the Euler number is with n = 57 terms of the defining series, since all digits (80) of a BigFloat value are exact:

```
EULER(56) #=
2.718281828459045235360287471352662497754969541622429154734
```

<sup>&</sup>lt;sup>3</sup> In mathematics and computer science, *currying* is the technique of translating the evaluation of a function that takes multiple arguments into evaluating a sequence of functions, each with a single argument.

```
483565013216246202514 =#

EULER(57) #=

2.718281828459045235360287471352662497754969541622429154734

483565013216246202549 =#

EULER(58) #= The 58th element of the Euler series

2.718281828459045235360287471352662497754969541622429154734

483565013216246202549 =#
```

# 2.4 Julia's package Plasm.jl

After several years of research about Linear Algebraic Representation (LAR) and algebraic operations [?, ?, ?, ?] with sparse matrices and solid models, a new Julia package for geometric programming, named Plasm.jl, was developed while writing this book. We aim to finish the software version 1.0 before the book is published.

The work has mainly consisted of porting to Julia the Pyplasm library [?] written in Python years ago. Of course, both are open-source and downloadable from the web<sup>4</sup>. Our software plan is to realize in version 1.0 a significant extension of the language related to computation of space arrangements [?] and Boolean solid algebras [?].

Of course, the reader is warmly invited to install the latest version of Julia and to download our package Plasm. jl on computational environment. The installation is not strictly necessary since web access will be available. Still, it is always helpful, while learning a new language, to have your own environment where you are free to do any experiment.

# Plasm.jl

The main file of the package is named as usual with the name Plasm.jl of the package itself. Its primary function is to create the run-time executable of Plasm.jl, by calling the external references (i.e., the exported functionality) taken from other packages and to include the Julia code of other package files, in our case fenvs.jl, hpc.jl, and viever.jl. Note that its name starts uppercase, according to the Julia's convention for packages.

# fenvs.jl

The fenvs.jl file, whose name stands for "functional environments" implements in Julia the majority of the small exciting programs developed for the GP4CAD book and includes many primitives for surface design with various

<sup>&</sup>lt;sup>4</sup> Citation of pyplasm and plasm.jl URLs.

methods. Other needed functions and functions of general use can be created directly by the reader if savant in computer graphics or CAD.

## hpc.jl

The primary data structure is the Hpc (Hierarchical Polyhedral Complex) [?], based on convex cells, and extended to allocate general polyhedral cells, i.e., polyhedra of any (low) dimension, connected but generally nonconvex and with interior holes. The hpc.jl file contains the developed design and implementation of the very general and multidimensional geometric data structure called Hpc [?], that allows to accommodate all the geometric and solid algorithms and tools developed in this book. As we show in the book, the Hpc structure is straightforward, versatile, and general. It does not use any of the large number of highly specialized and very complex data structures invented for solid modeling. We will show that our topological approach to geometric computations only needs the linear algebra of sparse matrices and vectors.

# viever.jl

Finally, the viewer.jl file is used as the home of an interactive geometry viewer, used to interact with the shapes generated by Plasm codes either on the terminal screen of a laptop or on the HTML interface of a web browser program. This web viewer version was developed to view the Plasm models on the web and to write rich text examples and exercises within the pages of a web notebook, and hence without the need to install any software.

# 2.5 Julia REPL (Read-Eval-Print-Loop)

An interactive language shell is an interactive computer programming environment in a single terminal that takes user inputs from keyboard or file, executes them, and prints the result to the display. A program developed in REPL terminal environment (CLI – command language interface) is written and executed piecewise. This approach requires that the language executable code may work as an interpreter, i.e. combining translation of source lines in machine code with immediate execution.

## REPL editor

Julia comes with a full-featured interactive command-line REPL (read-eval-print loop) built into the julia executable [?]. Once installed, to enter Julia, open a terminal and digit "julia" after the system prompt "\$". The acknowledgement in Figure 2.3 will appear on the screen. Once the REPL starts, you

will be at the Julia prompt. The Julia REPL can operate in different prompt modes:

- Julia mode (the default),
- help mode,
- · Pkg mode, and
- shell mode.

To enter help, Pkg, or shell mode, place the cursor at the beginning of the Julia mode prompt and type a question mark (?), a closing bracket (]), or a semicolon (;), respectively.

The help mode is used to get information about the meaning or use (arguments, examples, etc.) of a function, as written by the developer in the function "docstring". The Pkg (package) mode covers many things, including managing package installations, developing packages, working with package registries and more. The shell mode allows the terminal user to run shell commands from the Julia REPL.

To return to Julia mode, place the cursor at the beginning of the prompt and press Backspace [?]. A simple and very useful interactive blog to start use REPL is https://blog.glcs.io/julia-replheading-starting-the-julia-repl.

Fig. 2.3 To run Julia by invoking its name on the commend-line of the terminal on computer screen. Note the "using Plasm" after the REPL prompt "> julia".

# OhMyREPL package

The Julia package **OhMyREPL** hooks into the Julia REPL and gives it many new features [?]. It allows for several enhancements, including: syntax highlighting; bracket highlighting; bracket completion; rainbow brackets; markdown syntax highlighting; fuzzy REPL history search.

### Julia REPL workflow

The most basic Julia workflows involve using a text editor in conjunction with the julia command line. A common pattern includes the following elements:

1. Put code under development in a temporary module. Create a file, say Tmp.jl, and include within it

```
module Tmp
<your definitions here>
end
```

2. Put your test code in another file. Create another file, say tst.jl, which begins with

```
import Tmp
```

and includes tests for the contents of Tmp. The value of using <code>import</code> versus <code>using</code> is that you can call reload("Tmp") instead of having to restart the REPL when your definitions change. Of course, the cost is the need to prepend Tmp. to uses of names defined in your module. (You can lower that cost by keeping your module name short.) Alternatively, you can wrap the contents of your test file in a module, as

```
module Tst
    using Tmp
    <scratch work>
end
```

3. The advantage is that you can now do using Tmp in your test code and can therefore avoid prepending Tmp. everywhere. The disadvantage is that code can no longer be selectively copied to the REPL without some tweaking.

Lather. Rinse. Repeat. Explore ideas at the julia command prompt. Save good ideas in tst.jl. Occasionally restart the REPL, issuing

```
include("Tmp.jl")
include("tst.jl")
```

Julia's REPL provides rich functionality that facilitates an efficient interactive workflow [?].

# 2.6 Geometric Programming examples

Here we show a couple of simple examples to show the compactness and expressive power of our geometric language.

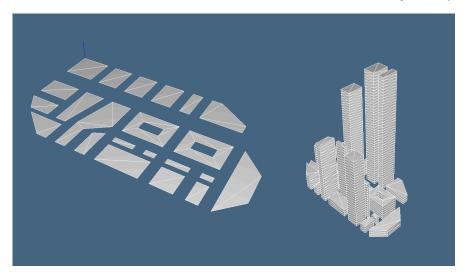


Fig. 2.4 (a) Hierarchical polyhedral complex (Lar) 2D value embedded in 3D. A perspective projection generates the image. Each connected shape is a polyhedral cell, even nonconvex and with inner holes. Each triangle is a convex cell; (b) 3D Plasm view of the model value of Hpc type discussed in Example 2.6.2

Coding 2.6.1 (2D virtual Manhattan) Of course, we start our geometric modeling session by telling the Julia compiler<sup>5</sup> to use the Plasm package. Then, we give the compiler the code and data that follows, possibly contained in a file Manhattan2D.jl.

```
julia> using Plasm
```

We start by defining the 2D coordinates of the vertices of our model.

```
julia> verts =
        [[0.,0],[3,0],[5,0],[7,0],[8,0],[9.5,1],[10,1.5],[0,3],
        [3,3],[5,3],[7,3],[8,3],[9.5,3],[0,4],[3,4],[5,4],[9.5,4],
        [12,4],[9.5,5],[10,5],[12,5],[0,6],[3,6],[5,6],[0,7],[3,7],
        [5,7],[9.5,7],[12,7],[9.5,8],[12,8],[0,9],[3,9],[5,9],[8,
        9],[9,9],[12,9],[0,10],[3,10],[5,10],[8,10],[9,10],[9.5,10],
        [10,10],[12,10],[6,11],[7,11],[0,12],[3,12],[9,12],[9.5,12],
        [0,13],[3,13],[6,13],[7,13],[9,13],[9.5,13],[0,14],[3,14],[5,
        14],[8,14],[9,14],[9.5,14],[10,14],[12,14],[0,15],[3,15],[5,
        15],[8,15],[0,16],[6,16],[7,16],[9,17],[9.5,17],[10,17],[12,
        17],[6,18],[7,18],[9,18],[9.5,18],[10,18],[12,18],[2,19],[3,
        19],[5,19],[8,19],[9,19],[9.5,19],[10,19],[12,19],[5,20],[12,
        20],[7,22],[10,22],[9,6],[12,6],[9,15],[9.5,15],[10,15],[12,
        15]]
```

<sup>&</sup>lt;sup>5</sup> To load Plasm in Julia, open a terminal, start your julia application, and after the prompt julia> write using Pkg; Pkg.add("Plasm")

Then we give the following description of convex cells of type Cells = Vector {Vector{Int64}}:

```
julia> cells =
    [[1,2,9,8],[3,4,11,10],[5,6,13,12],[14,15,23,22],[16,
17,19,24],[7,18,21,20],[25,26,33,32],[27,95,28,35,34],[95,
96,29,28],[30,31,37,36],[38,39,49,48],[40,41,47,46],[41,61,
55,47],[55,61,60,54],[54,60,40,46],[42,43,51,50],[44,45,65,
64],[52,53,59,58],[56,57,63,62],[66,67,84,83,70],[68,69,72,
71],[69,86,78,72],[78,86,85,77],[71,77,85,68],[97,98,74,
73],[99,100,76,75],[79,80,88,87],[81,82,90,89],[91,92,94,93]]
```

Finally, verts and cells are transformed in a geometric object of type Hpc by the function MKPOL and stored in the Julia variable named model.

```
julia> model = MKPOL(verts,cells)
# Hpc ... ...
```

A model image is generated by the Plasm viewer, within a system window named "Manhattan2D", for interaction with mouse and arrow buttons.

```
julia> VIEW( model, "Manhattan2D" )
```

From terminal we might write: \$ julia path/Manhattan2D.jl

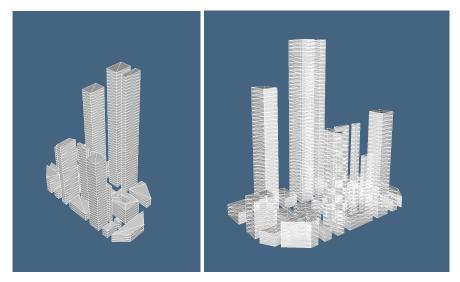


Fig. 2.5 3D model generated by using the 2D cells defined in Manhattan2D example.
(a) opaque visualization; (b) transparent visualization from a different viewpoint.

Coding 2.6.2 (3D virtual Manhattan) The model of Figure 2.5 is given by (1) a vector ManhattanH of floor numbers; (2) generation of corresponding 1D and 2D vectors of Hpc objects stored in pols1D and pols2D; (3) by Cartesian product of corresponding Hpc pairs (2D,1D). See below:

```
pols3D = AA(splat(*))(TRANS([pols2D, pols1D]))
# 29-element Vector{Hpc}:
VIEW(STRUCT(pols3D), "Manhattan3D")
```

In particular, we define an array of virtual heights for each *polygon* of Manhattan2D. Such numbers are transformed in repeated storeys heights by AA(DIESIS) (where DIESIS was the # operator in FL-based Plasm, but it is not usable in Julia, since it denotes comments) and then codified as 1D Hpc polyhedra stored in pols1D by the QUOTE operator. Analogously, a set of 2D polyhedra is stored in the pols2D array. The Cartesian product \* of corresponding Hpc objects is stored in pols3D array using the Julia's splat function. Finally, a single Hpc object is visualized in a system window named Manhattan3D.

It is worth noting that every polygon\*segment multiplication of two Hpc objects produces a new Hpc polyhedron of dimension equal to the sum of dimensions of operands. The reader should not forget that the Plasm language and the Hpc data structure are both *multidimensional*.

We also note that a large number of significant programming examples with Julia and Plasm in Julia can be found inside the file Plasm/fenvs.jl, and executed in the terminal by writing \$ julia ./test/fenvs.jl, being located into the Plasm.jl directory, and after having downloaded and installed the package Plasm.jl in a recent julia environment.

Coding 2.6.3 (Solid 2D graph of a scalar function) A simple unusual geometric example is given here, showing some of canonical constructs of geometric design with Plasm. First, we build and show in Figure 2.6a the Domain2D of the parametric 2D solid model given in Figure 2.6b. The MAP operator is applied first to the function Mapping2D to be applied to all vertices of a cell decomposition of Domain2D, in turn generated as the Cartesian product POWER of two 1D cellular complexes, so producing a 2D cell complex of the mapped interval Domain2D with  $36 \times 6$  squared cells.

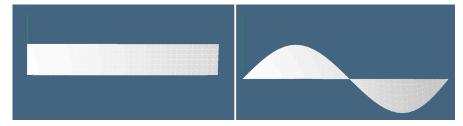


Fig. 2.6 The mapping of a function of two arguments on its 2D parameter domain. Each element (point and cell) of the cellular domain is paired with a corresponding element in the function range. The figure shows how the elements are paired. (a) The cellular complex decomposing the function domain  $[2\pi, 1]$ ; (b) the function range transformed after the function  $\mathbf{p} \rightarrow [\mathbf{u}, \sin(\mathbf{u}) * \mathbf{v}]$  was mapped on the vertices of the 2D domain.

```
Domain2D = Power(INTERVALS(2π)(36), INTERVALS(1)(6));
Mapping2D = p->((u,v)=p; [u,sin(u)*v]);
model = MAP(Mapping2D)(Domain2D)
```

The two parameters u, v of the model domain are worked out by p->(...), anonymous Julia function, and applied to each vertex of Domain2D by the MAP operator. Two geometric objects Domain2D and model are finally shown and displayed in Figure 2.6:

```
VIEW(Domain2D)
VIEW(model)
```

The virtual models of both Domain2D and model, finally visualized in Figure 2.6, are made available for display by the Plasm operator VIEW, and can be interactively handled by the user.

# Chapter 3 Geometry and topology primer

It is implausible to set up a computer language and computational environment to define and deliver the geometric shape of construction elements and spaces without a pretty good understanding of basic geometric and topological concepts. Hence, this chapter introduces some basic notions about geometric spaces, operators, and properties, like linearity, that provide the computational foundation of shape definition and model construction. The essential elements of geometric spaces and cellular models, like affine transformations and simplicial, cubical, and polyhedral complexes, constitute the substrate of Plasm discourse for building hierarchical assemblies of AEC products of any scale and complexity.

# 3.1 Geometric Spaces

In mathematics and science, a space is a set of objects with some added structure. In this section we discuss the geometrical spaces used to represent inside a computer memory the actual objects of natural, artificial or virtual environments, or better their geometric models, in order to study, visualize and/or simulate some associated physical behavior. In particular, we deal in this section with linear, affine, and convex spaces as sets of vectors and/or points with selected properties.

# 3.1.1 Vector space

The concept of vector space is undoubtedly the most helpful instrument of mind invented by mathematicians for scientists, engineers, and architects, to represent and study formally or graphically both the primary and complex arrangements and behaviors of our natural and artificial environment.

#### Definition

A vector space (or linear space)  $\mathcal{V}$  over a field  $\mathcal{F}$  is a set, closed w.r.t. two composition rules (the output of composition belongs to the set of inputs).

The elements  $v \in \mathcal{V}$  are called *vectors*, and are often represented by an oriented arrow with given direction, orientation, and length. The elements  $\alpha \in \mathcal{F}$  are called *scalars*. The sum of two non-zero vectors  $u, v \in \mathcal{V}$  is a third vector  $w = u + v \in \mathcal{V}$  with direction and length different from both u and v. We may write v + v = 2v, and see here both the operations of a vector space: (a) addition of vectors, and (b) multiplication times a scalar.

The product of a vector by a scalar  $\alpha \mathbf{v} = \mathbf{v} \alpha \in \mathcal{V}$  is a vector collinear with  $\mathbf{v}$  and with different length if  $\alpha \neq 1$ . This explain the name "scalar" since a number "scales" (change the length of) the vector it multiplies.

The length of  $u = \alpha v$  grows or shrinks w.r.t. v according to a positive  $0 < \alpha < 1$ . If  $\alpha < 0$ , then u has orientation opposite to v.

## Linear independence

A linear combination of vectors is a new vector that is defined as a sum of scalar multiples of other vectors. Let  $v_1, v_2, ..., v_n \in \mathcal{V}$  and  $\alpha_1, \alpha_2, ..., \alpha_n \in \mathcal{F}$ , with  $\mathcal{V}$  a vector space on the field  $\mathcal{F}$  of scalar numbers. The vector

$$\boldsymbol{w} = \alpha_1 \boldsymbol{v}_1 + \alpha_2 \boldsymbol{v}_2 + \dots + \alpha_n \boldsymbol{v}_n = \sum_{i=1}^n \alpha_i \boldsymbol{v}_i \in \mathcal{V}$$

is called a linear combination of vectors  $v_1, v_2, ..., v_n$ .

- Two or more vectors are said *linearly independent* if none of them can be written as a linear combination of the others;
- if at least one of them can be written as a linear combination of the others, then they are said *linearly dependent*.

Given a set of vectors, you can determine if they are linearly independent by writing the vectors as the columns of a matrix A, and solving Ax = 0.

If there are non-zero solutions, then the vectors are linearly dependent. If the only solution is x = 0, then they are linearly independent.

The typical vector spaces in this book will be (a) the numeric field  $\mathbb{R}$  for scalars and  $\mathbb{R}^n$  for vector coordinates, with (b) dim = d, for  $0 \le d \le 3$ .

### Examples

Coding 3.1.1 (Matrices  $m \times n$  are vector spaces). Generate a random  $100 \times 100$  matrix A = rand(100,100). The default of rand() function are values of type Float64 in [0.0,1.0], so we ask the compiler to compute

the matrix multiplication by the scalar  $\pi$ , denoted by the Greek symbol. The generated value is shown commented and simplyfied for printing in small space:

```
A = rand(100,100) * π #=

100x100 Matrix{Float64}:

0.901069 ... 0.172854

0.357548 0.8537

...

0.519136 0.857736 =#
```

Coding 3.1.2 (Random vector generation.) Then, we generate a 100-vector of random integers within the interval [0,100], by using the Julia 0:100 iterator:

```
b = rand(0:100, 100) #=
100-element Vector{Int64}:
61
85
:
51 =#
```

Coding 3.1.3 (Multiplication matrix-vector.) This operation is implemented natively and very efficiently for big matrices in Julia, as well the product and the sum of dense matrices (the detail of printing depends on available space on output screen):

```
A * b #=
100-element Vector{Any}:
2786.383421064984
2554.1378635659034
:
2735.1205994270754 =#
```

Coding 3.1.4 (Addition matrix-vector.) Conversely, the sum of a matrix with a vector is not a linear operation, so we need to broadcast (. operator) the vector b on all columns of A:

Summing up: while the Matrix by Vector multiplication is a native operation on the linear space of Numbers, The addition of Matrix and Vector is not, and the broadcast operator must be used. Same for the sum of a matrix and a single scalar value. The reader should try.

### Subspace

Let  $\mathcal{V}$  be a vector space on the field  $\mathcal{F}$ . We say that  $\mathcal{U} \subset \mathcal{V}$  is a *subspace* of  $\mathcal{V}$  if  $\mathcal{U}$  is a vector space with respect to the same operations. In particular,  $\mathcal{U} \subset \mathcal{V}$  is a *subspace* of  $\mathcal{V}$  if and only if:

- 1.  $\mathcal{U} \neq \emptyset$ ;
- 2. for each  $\alpha \in \mathcal{F}$  and  $\mathbf{u}_1, \mathbf{u}_2 \in \mathcal{U}$ ,  $\alpha \mathbf{u}_1 + \mathbf{u}_2 \in \mathcal{U}$

The *codimension* of a subspace  $\mathcal{U} \subset \mathcal{V}$  is defined as dim  $\mathcal{V} - \dim \mathcal{U}$ .

It may be useful to note that the intersection of subspaces is a subspace. In particular, if  $\mathcal{U}_1, \mathcal{U}_2$  are subspaces of  $\mathcal{V}$ , then  $\mathcal{U}_1 \cap \mathcal{U}_2$  is a subspace of  $\mathcal{V}$ .

## Generators, Bases, and Coordinates

Span and generators

The smallest subset of vectors that can be generated by linear combinations of a subset  $S \subset \mathcal{V}$  of linearly independent vectors is called  $\operatorname{span} S$ .

The set S is therefore called a set of *generators*.

The span S is closed with respect to addition and multiplication, and hence is a subspace including the zero vector, which is contained in every subspace.

### Bases and coordinates

The *basis* and *dimension* of the linear space  $\mathcal{V}$  are a minimum set of generators for  $\mathcal{V}$ , and its number  $d = \dim \mathcal{V}$  of elements, respectively.

Every basis of a linear space  $\mathcal{V}$  has the same number d of elements.

When a basis for  $\mathcal V$  has been fixed, i.e. an ordered minimal subset of generators  $B\subset \mathcal V$  has been chosen, every vector  $\boldsymbol v\in \mathcal V$  can be expressed uniquely as the linear combination of elements of B with scalars. The ordered tuple of such scalars is called the *coordinate* tuple, or the coordinates, of vector  $\boldsymbol v\in \mathcal V$ , and denoted as  $[\boldsymbol v]$ .

Remark 3.1 It is to mention that to represent vectors by coordinates requires that a minimum set of space generators and their ordering have been already chosen. We will say that the space has been parameterized, since every vector is uniquely identified by the linear combination of the basis elements with a unique tuple of scalars.  $\Box$ 

Remark 3.2 The basis is often denoted by the ordered sequence  $(e_1, \ldots, e_d)$  of vector elements, or by the matrix  $[e_1 \cdots e_d]$  of their coordinates by columns, where  $e_i = [0, 0, 1, \ldots, 0]^t$  is a (column) tuple of zeroes, with only one element 1 in position *i*. The *standard basis* of a coordinate vector space is the set of vectors whose components are all zero, except one that equals 1.

# Examples of vector spaces

The usual geometric example of vector space has the oriented arrows<sup>1</sup> as elements, summed with the parallelogram rule, and scaling related to elongation or shortening. Other examples are the linear spaces of matrices  $\mathcal{M}_m^n(\mathbb{R})$  with real elements, m rows and n columns, and  $\mathcal{M}_m(\mathbb{R})$  and  $\mathcal{M}^n(\mathbb{R})$  of column and row vectors, respectively.

Linear space is also the space  $\mathbb{P}_n(\mathbb{R})$  of real polynomial functions  $p: \mathbb{R} \to \mathbb{R}$  such that  $x \mapsto p^n(x)$  of degree  $\leq n$ . In particular,  $p^n(x) = a_0 + a_1 x + a_2 x^2 + \cdots + a_{n-1} x^{n-1} + a_n x^n$  is exactly a linear combination of a tuple of n+1 scalars  $a_k$ ,  $0 \leq k \leq n$ , with the *power basis* of polynomials  $x^k$ ,  $0 \leq k \leq n$ .

# The Bernstein bases of polynomial space $\mathbb{P}_n(\mathbb{R})$

In geometric modeling and computer graphics, the Bernstein-Bézier basis of polynomials is fundamental. The n+1 Bernstein polynomials of degree n, defined as

$$B_k^n(x) = \binom{n}{k} x^k (1-x)^{n-k}, \qquad 0 \le k \le n$$

form a basis for the vector space  $\mathbb{P}_n(\mathbb{R})$  of polynomials of degree at most n with real coefficients. Therefore, to implement the basis and to draw a graph of each basis function we have to consider: the degree n we are interested to; the ordinal number k of each function  $(1 \leq k \leq n)$ ; and finally the independent variable x such that  $x \mapsto p^n(x)$ .

## Script 3.1.1 (Pure functional style in Julia)

The julia function B given here returns the whole range of Bézier-Bernstein polynomials of any degree, and is very useful in curve geometric modeling.

$$B = n -> k -> u -> binomial(n,k) * u^k * (1-u)^(n-k)$$

Let us make some checks: B(1)(0)(0.5) == B(1)(1)(0.5) == 0.5 #=> true, for the degree-1 basis made by two polynomials B(1)(0) and B(1)(1). The vhole basis of degree n is generated by the higher-level function

<sup>&</sup>lt;sup>1</sup> Formally: the equivalence classes of equipollent oriented arrows. In Euclidean geometry, equipollence is a binary relation between directed line segments.

Bernstein(n) that we test in the quadratic case, where it returns a 3—element array of Vector{Function}:

Coding 3.1.5 (Generating Bernstein polynomial bases) The function Bernstein(n) is defined by mapping the partial function B(n) over the integer array [0, 1, ..., n].

Now, we go to generate a discrete sequence of functions for the "vector" function Bernstein(2), in order to test the implementation. Above we see also the second function of Bernstein basis of degree n=2, and finally its value 0.18 computed for x=0.1.

Coding 3.1.6 (Sampling of third quadratic polynomial) The function Bernstein(2)[3] denotes the third function of the vector array Bernstein(2) of type ::  $Vector{Function}$ , and we compute the three function values for x=0.5, that you can check in Figure 3.1.

```
Bernstein(2)[3] #=
37 (generic function with 1 method) =#

Bernstein(2)[1](0.5) # => 0.25
Bernstein(2)[2](0.5) # => 0.5
Bernstein(2)[3](0.5) # => 0.25
```

Coding 3.1.7 (Sampling of third basis polynomial of degree 4) . We may look at the sequence of pairs x, Bernstein(2)[3](x). A sampling of x values is created by collect(0:0.1:1):

```
m = 10
for u in collect(0:1/m:1)
    println(" $(u), $(Bernstein(4)[3](u)) ")
end      #=
0.0, 0.0
0.01, 0.0005880600000000001
0.02, 0.00230496
0.03, 0.00508086
: : =#
```

The Bernstein basis enjoy interesting properties. They are  $\geq 0$  for every value of the independent variable  $x \in [0,1]$ . Even more, for every x the elements of each basis sum to 1. Such bases are said partition of unity.

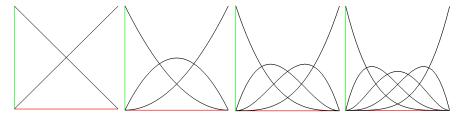


Fig. 3.1 The four images give the graph, in  $[0,1] \times [0,1] \subset \mathbb{E}^2$ , of Bernstein's basis of linear, quadratic, cubic, and quartic polynomials in linear spaces  $\mathbb{P}_n(\mathbb{R})$ , with  $1 \leq n \leq 4$ , and dim $(\mathbb{P}_n(\mathbb{R})) = n + 1 = 2, 3, 4$ , and 5, respectively.

Coding 3.1.8 (Example of FL & Plasm Combinators) We remark on the incredible expressive power of FL programming inherited in Plasm.jl.

It is greatly useful to analyze goals and types of subexpressions, recuperating backward the development process to write the result expression. First, we define the 1D model dom1D of the function domain [0, 1] with 36 intervals, which is a geometric value of Hpc type:

Then, consider the  $\mathbb{E} \to \mathbb{E}^2$  vector function CONS(...) $\circ$ S1 of one variable, that is applied to the vertices of dom1D::Hpc by the MAP operator.

The selector function S1 is used to extract the first (and unique) coordinate value from the internal data structure point array. The CONS operator, acting on argument of Vector{Function} type, transforms a function array into a vectorial function of coordinate functions x(u), y(u), etc.

Hence, we have  $F = CONS(::Vector{Function}) \circ Sd : \mathbb{E}^d \to \mathbb{E}^n$ , where d is the number of coordinates of domain vertices (one in this case), and n is the dimension of the embedding space, i.e., the number of coordinate functions inside the array argument, i.e. CONS([,]), two in this case:

```
F(k) = CONS([ID, Bernstein(4)[k]]) oS1 #= #122 (generic function with 1 method) =#
```

Finally, we note that expressions like MAP(F)(dom1D) are used to apply the F vector function to an object of Hpc type, normally to bend it. Remember that we generated an array of Hpc curves, for k=1:n+1 — in our case with n=4. For this purpose, we need to finally apply the STRUCT operator, to transform the array of curves into a single Hpc value.

STRUCT( MAP(F(k))(dom1D) for k=1:n+1 )

## Change of basis

It is often necessary, given a basis in a linear space, to compute a novel parameterization of the space with respect to a different, possibly more simple or useful basis. In other words, may be necessary to compute new coordinates for the vectors of the space, with respect to a new basis. This operation is called *change of basis*.

For concreteness, let we have vector data in a 3D linear space, where we need to compute their representation in a different basis. Let us denote the new basis as  $(u_i)$ , and the standard one as  $(e_i)$ , with  $i \in \{1, 2, 3\}$ . We may write the change of coordinates as a matrix map  $T : \mathcal{V} \to \mathcal{V}$ , such that  $[u_{ij}] \mapsto [e_{ij}]$ , so transforming the old coordinates of the new basis into the standard basis.

Therefore we set  $T[u_{ij}] = [e_{ij}]$  and, since it is  $[e_{ij}] = \mathbb{I}_3$ , i.e., the  $3 \times 3$  identity matrix, the solution is  $T = [u_{ij}]^{-1}$ . Of course, this matrix exists provided that  $\det[u_{ij}] \neq 0$ . In other words, the vectors of the new basis must be linearly independent.

## 3.1.2 Affine space

In geometric modeling and computer graphics it is useful to distinguish between a space of vectors and a space of points (supported by vectors). A space of points, that provides for an operation of *displacement*, is called an *affine space*, and is represented here by the  $\mathcal{A}$  symbol. In particular, we call affine action the function:

$$\mathcal{A} \times \mathcal{V} \to \mathcal{A}$$
,

so that the *displacement* from a point  $a \in \mathcal{A}$  to a point  $b \in \mathcal{A}$  is given. An affine space  $\mathcal{A}$  is endowed with an operation of *difference* of points  $\mathcal{A} \times \mathcal{A} \to \mathcal{V}$  where

$$a + v = b$$
, and hence  $b - a = v \in \mathcal{V}$ 

### Remarks

We note that a **vector space** provides for an internal operation of (a) *sum* of two vectors and the external (b) *product* of a vector by a scalar, both returning a vector. Conversely, in an **affine space** we have an external operation of

(i) difference of points, returning a vector, and a (ii) sum of a point and a vector, called displacement, returning a point.

Even more, in a vector space  $\mathcal{V}$  there is a distinguished element 0, the zero vector, which is contained in all subspaces. On the contrary, in an affine space  $\mathcal{A}$  all points are equivalent, with no distinguished elements.

We usually say that a space of points has been parameterized when a *Cartesian system* (origin and basis) has been chosen. In order to use coordinates to make it easier to work with points we have to choose:

- 1. a point, called *origin*, to associate with the zero vector, and
- 2. an orthonormal basis of vectors.

The above steps consent to associate each point with the tuple of coordinates of its displacement vector from the origin.

In a vector space all the subspaces have at least a common element, the zero vector. Contrariwise, two affine subspaces may not have common elements. In such a case they are said *parallel*.

The dimension n of an affine space  $\mathcal{A}$  is that of its supporting vector space  $\mathcal{V}$ . A common word for affine subspaces of d dimension is d-hyperplane, or d-hyperspace when d > 3. Lines and planes are affine subspaces of dimension 1 and 2, respectively.

## Affine independence and local parameterization

In an affine space  $\mathcal{A}$  of sufficiently high dimension n we say that two points are affinely independent when non coincident, three points when non aligned, four points when non coplanar, and so on.

In general, d+1 points  $(d \leq n)$  are affinely independent when the d vectors defined by the differences  $\mathbf{p}_k - \mathbf{p}_0$  of points from one of them are linearly independent.

The affine independence of a subset of d+1 points is often used to establish local coordinate systems on lines, planes and higher dimensional subsets of points. Let us choose two non coincident points a, b on a 3D line. Any other point p of the line remains parameterized by the  $\alpha$  scalar in the expression

$$p = a + \alpha(b - a).$$

Note that b - a is a vector,  $\alpha(b - a)$  is a vector,  $a + \alpha(b - a)$  is a point plus a vector, which is a point. The typing of our expression looks correct!

Analogously, let us choose three non colinear points a, b, c in 3D space. They are certainly non coincident, and of course fix a plane, i.e., an unique affine subspace of dimension 2 embedded in space. A parameterization of this plane is given by the pairs  $(\alpha, \beta)$  of scalars in the expression

$$q = \alpha(b - a) + \beta(c - a),$$

as the reader may immediately check. Similar local coordinates will hold in every affine d-subspace in n-dimensional space.

The typical affine space of points used in this book is the Euclidean space  $\mathbb{E}^d$ ,  $1 \leq d \leq 3$ , usually furnished of a Cartesian system, with coordinates in Float64.

# 3.1.3 Convex space

Let us consider the Euclidean space  $\mathbb{E}^d$ ,  $d \in \{2,3\}$  affine over the reals, that is the fundamental space of geometry, intended to represent physical space.

Affine subspaces become *convex sets* when a numeric constraint is imposed on the possible parameter values of an affine combination of points. Two points  $\boldsymbol{a}, \boldsymbol{b} \in \mathcal{A}$  become the extreme elements of a line segment  $\boldsymbol{p} = \alpha \, \boldsymbol{a} + (1 - \alpha) \, \boldsymbol{b}$  as set of points, by adding the further constraint  $\alpha + \beta = 1$  to the parameter values  $\alpha, \beta \geq 0$ , so posing  $\beta = 1 - \alpha$ . Analogously, setting  $\alpha + \beta + \gamma = 1$  and  $\alpha, \beta, \gamma \geq 0$  constraints the set of points combination of three non aligned points  $(\boldsymbol{a}, \boldsymbol{b}, \boldsymbol{c})$ .

### Positive, Affine and Convex Combination of Points

Let  $p_1, p_2, \ldots, p_n$  be affinely independent points in  $\mathbb{E}^d$ , and  $\alpha_1, \alpha_2, \ldots, \alpha_n$  be scalar in  $\mathbb{R}$ . Their combination  $\alpha_1 p_1 + \alpha_2 p_2 + \cdots + \alpha_n p_n$  is said *positive*, affine, and *convex*, respectively, when

1. 
$$\alpha_1, \alpha_2, \dots, \alpha_n \ge 0$$
 (positive combination)  
2.  $\alpha_1 + \alpha_2 + \dots + \alpha_n = 1$  (affine combination)

3. 
$$\alpha_1 + \alpha_2 + \cdots + \alpha_n = 1$$
 and  $\alpha_1, \alpha_2, \dots, \alpha_n \ge 0$  (convex combination)

It may be interesting to verify that the affine combination of points is a point. Let us eliminate the  $\alpha_1$  scalar using the unitary sum of scalars constraint:

$$\mathbf{p} = (1 - \alpha_2 - \dots - \alpha_n)\mathbf{p}_1 + \alpha_2\mathbf{p}_2 + \dots + \alpha_n\mathbf{p}_n$$

$$= \mathbf{p}_1 + \alpha_2(\mathbf{p}_2 - \mathbf{p}_1) + \dots + \alpha_n(\mathbf{p}_n - \mathbf{p}_1)$$
(3.1)

which, of course, is a point in  $\mathbb{E}^d$ .

A convex combination is positive and affine.

The set of all convex combinations of points  $C \subset \mathbb{E}^n$  is the *convex hull* of C. The convex hull is the smallest compact set containing the points in C. It is the intersection of all compact sets of  $\mathbb{E}^d$  which contain hull C.

Convex coordinates of a point  $c \in \text{hull } C$  are the scalars whose convex combination with C elements produces c. If  $C \subset E^n$  has n+1 affinely inde-

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pendent elements, i.e., it is a simplex (see Section ??) the convex coordinates of each  $c \in \text{hull } C$  are unique.

# Characterization of affine pyramids

Speaking of local parameterization of affine subspaces, we did not fix any constraint on the signs of the scalar parameters (usually in the field  $\mathbb{R}$  of reals). If all the scalars are positive, then all the spanned points stay inside the interior of a planar (or solid) angle contained within the lower-dimensional affine subspaces generated. It would be not difficult to see that the whole subspace will be partitioned by an arrangement of subspaces centered on the fixed point of the set of linearly independent vectors. As an example, consider the partitioning of 3D space generated by the  $8=2^3$  octants defined by 4 non coplanar points, and in particular those generated when three points are at unit distance by the first, and the three difference vectors are pairwise orthogonal.

### Positive, Affine and Convex coordinates

Affine Coordinates for Convex Sets
Barycentric Coordinates for Convex Sets
Test for boundary points: interior and exterior

## 3.2 Cellular models

This section discusses computational topology concepts used to compute the 2D/3D space partition induced by a collection of geometric objects of dimension 1D/2D, respectively. Cellular models are are often called *meshes* in engineering and design jargon. A mesh is a representation of a domain by discrete cells. In many areas of geometric/numeric computational engineering, including geo-mapping, computer vision and graphics, finite element analysis, medical imaging, geometric design, and solid modeling, one has to compute incidences, adjacencies and ordering of mesh cells, generally using disparate and incompatible data structures and algorithms. Only sparse vectors and matrices are used in Plasm to compute the linear spaces of chains, called chain complex, from dimension zero to three.

# Topological definitions

A complex is a graded set  $S = \{S_i\}_{i \in I}$  i.e. a family of sets, indexed here over  $I = \{0, 1, 2, 3\}$ . We use two different but intertwined types of complexes, and specifically complexes of *cells* and complexes of *chains*. Their definitions and some related concepts are given in this section. Greek letters are used for the cells of a space partition, and roman letters for chains of cells, coded as either (un)signed integers or sparse arrays of (un)signed integers.

**Definition 3.1** (d-Manifold) A manifold is a topological space that resembles a flat space locally, *i.e.*, near every point. Each point of a d-dimensional manifold has a neighborhood that is homeomorphic<sup>2</sup> to  $\mathbb{E}^d$ , the Euclidean space of dimension d. Hence, this geometric object is often referred to as d-manifold.

Homeomorphic neighborhood means "topologically equivalent", like a rubber patch, that can be stretched without changing its topology.

**Definition 3.2 (Cell)** A p-cell  $\sigma$  is a p-manifold with boundary  $(0 \le p \le d)$  which is piecewise-linear, connected, possibly non convex, and not necessarily contractible. This definition refers to cellular complexes used in this paper and is different from other ones because a cell is neither simplicial, nor convex, nor contractible.

In Plasm, cells may contain internal holes; cells of CW-complexes [?] are, conversely, contractible to a point. We deal with Piecewise-Linear (PL) cells of dimensions 0, 1, 2, and 3, respectively. It should be noted that 2- and 3-cells may contain holes, while remaining connected. In other words, Plasm cells are p-polyhedra, i.e. segments, polygons and polyhedrons embedded in 2D or 3D space.

**Definition 3.3 (Cellular complex)** A cellular p-complex is a finite set of cells that have at most dimension p, together with all their r-dimensional boundary faces  $(0 \le r \le p)$ . A face is an element of the PL boundary of a cell, that satisfy the boundary compatibility condition. Two p-cells  $\alpha, \beta$  are said boundary-compatible when their point-set intersection contains the same r-faces  $(0 \le r \le p)$  for both  $\alpha$  and  $\beta$ .

A cellular *p*-complex is said *homogeneous* when each *r*-cell  $(0 \le r \le p)$  is face of a *p*-cell.

**Definition 3.4 (Skeleton)** The s-skeleton of a p-complex  $\Lambda_p$  ( $s \leq p$ ) is the set  $\Lambda_s \subseteq \Lambda_p$  of all r-cells ( $r \leq s$ ) of  $\Lambda_p$ . Every skeleton of a regular complex is a regular subcomplex. The difference  $\Lambda_r - \Lambda_{r-1}$  of two skeletons is the set  $U_r$  of r-cells.

<sup>&</sup>lt;sup>2</sup> Homeomorphic means opologically equivalent.

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**Definition 3.5 (Support space)** The support space  $\Lambda$  of a cellular complex is the point-set union of its cells.

**Definition 3.6 (Characteristic function)** Given a subset S of a larger set A, the characteristic function  $\chi_A(S)$ , also called the *indicator function*, is the function defined to be identically one on S, and zero elsewhere. [?].

# 3.2.1 Simplicial complex

**Definition 3.7 (Join operation.)** The *join* of two compact sets of points  $P, Q \subset \mathbb{E}^n$  is the set PQ of convex combinations of points in P and in Q. The join operation is associative and commutative.

**Definition 3.8 (Simplex.)** A *d-simplex*  $\sigma_d \subset \mathbb{E}^n$   $(0 \le d \le n)$  may be defined as the repeated join of d+1 affinely independent points, called *vertices*.

A d-simplex can be seen as a d-dimensional triangle: a 0-simplex is a point, a 1-simplex is a segment, a 2-simplex is a tetrahedron, and so on.

Coding 3.2.1 (Plasm d-dimensional simplex) A d-simplex is generated by the Plasm package by the SIMPLEX multi-dimensional function:

```
julia> using Plasm
julia> SIMPLEX::Function
SIMPLEX (generic function with 1 method)
```

Coding 3.2.2 (Dataset associated to simplices.) The simplex datasets follow, for the first integer parameters. Note that SIMPLEX(d) contain d+1 coordinate points of dimension d, and that hulls fields contain d+1 indices of points, i.e. a single convex cell:

```
SIMPLEX(0) #=
Hpc(MatrixNd(1), Geometry([Float64[]], hulls=[[1]])) =#

SIMPLEX(1) #=
Hpc(MatrixNd(2), Geometry([[0.0], [1.0]], hulls=[[1, 2]])) =#

SIMPLEX(2) #=
Hpc(MatrixNd(3), Geometry([[0.0, 0.0], [1.0, 0.0], [0.0, 1.0]], hulls=[[1, 2, 3]])) =#

SIMPLEX(3) #=
Hpc(MatrixNd(4), Geometry([[0.0, 0.0, 0.0], [1.0, 0.0, 0.0], [0.0, 1.0]], hulls=[[1, 2, 3, 4]])) =#
```

```
SIMPLEX(4) #=
Hpc(MatrixNd(5), Geometry([[0.0, 0.0, 0.0, 0.0], [1.0, 0.0, 0.0, 0.0], [0.0, 1.0, 0.0, 0.0], [0.0, 0.0, 1.0, 0.0], [0.0, 0.0, 1.0, 0.0], [0.0, 0.0, 1.0], hulls=[[1, 2, 3, 4, 5]])) =#
```

For the sake of visual simplicity, we remove the julia> prompt for Plasm scripts, and use the multilinear comment to reduce their visual rumor.

**Definition 3.9** Skeleton and Faces. The set  $\{v_0, v_1, \ldots, v_d\}$  of vertices of  $\sigma_d$  is called the  $\theta$ -skeleton of  $\sigma_d$ . The s-simplex generated from any subset of s+1 vertices  $(0 \le s \le n)$  of  $\sigma_d$  is called an s-face of  $\sigma_d$ .

Remark 3.3 Let us notice, from the definition, that a simplex may be considered both as a purely *combinatorial object* and as a *geometric object*, i.e. as the compact point-set defined by the convex hull of a discrete set of points.  $\square$ 

**Definition 3.10 (Simplicial Complex)** A set  $\Sigma$  of simplices is called a *triangulation*. A *simplicial complex*, often simply denoted as *complex*, is a triangulation  $\Sigma$  that verifies the following conditions:

- 1. if  $\sigma \in \Sigma$ , then any face of  $\sigma$  belongs to  $\Sigma$ ;
- 2. if  $\sigma, \tau \in \Sigma$ , then either  $\sigma \cap \tau = \emptyset$ , or  $\sigma \cap \tau$  is a face of both  $\sigma$  and  $\tau$ .

A simplicial complex can be considered a *well-formed* triangulation. Such kind of triangulations are widely used in engineering analysis, e.g., in topography or in finite element methods.

Coding 3.2.3 (binomial numbers and simplex faces) There are

$$\sum_{k=0}^{n} \binom{n}{k} = 2^n$$

k-faces in a n-simplex. Testing this statement is simple and very elegant with Julia. We may verify, e.g., that simplex(5) faces are a set of  $2^5$  elements:

```
julia> [binomial(5,k) for k=0:5]'
1x6 adjoint(::Vector{Int64}) with eltype Int64:
1  5  10  10  5  1

julia> [binomial(5,k) for k=0:5] |> sum == 2^5
true
```

Coding 3.2.4 A simple coding example produces the simplicial complex providing the triangulation of the boundary of a convex hull.

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```
julia> p = rand(6,3)
                                     #=
6×3 Matrix{Float64}:
0.456121 0.340689
                     0.523394
           0.920846
 0.670731
 0.511325
           0.83709
                     0.765548
 0.27295
           0.344676
                     0.246891
 0.155611
                     0.372059
 0.997037
           0.689132
                     0.594624
                                       =#
```

The p matrix holds 6 random 3D points. The q matrix receives an array of arrays:

```
julia> q = [p[k,:] for k=1:size(p,1)] #=
6-element Vector{Vector{Float64}}:
  [0.4561213293752391, 0.34068894716553066, 0.5233939444809501]
  [0.6707310911896536, 0.9208461259235498, 0.8105811405026019]
  [0.5113250218604997, 0.8370897242704682, 0.7655476328700838]
  [0.2729499977250678, 0.3446764528053594, 0.24689130455109154]
  [0.15561059083290985, 0.26258788197490757, 0.37205895443742]
  [0.9970371401112009, 0.6891321979374976, 0.5946242627157257]
  =#
```

Their convex hull is computed by the Plasm function CONVEXHULL and stored as Hpc data structure:

And finally the dataset is transformed into a Lar data structure, which contains all the boundary polygons :FV and edges :EV.

We remark that the Hpc describes only the *convex cells* of its dataset. Conversely, the Lar data structure gives explicitly all cells of any dimension.  $\Box$ 

The order of a complex is the maximum order of its simplices. A complex  $\Sigma_d$  of order d is also called a d-complex. A d-complex is said to be regular or pure if each simplex is a face of a d-simplex. A regular d-complex is homogeneously d-dimensional.

The combinatorial boundary  $\Sigma_{d-1} = \partial \sigma_d$  of a simplex  $\sigma_d$  is a simplicial complex consisting of all proper s-faces (s < d) of  $\sigma_d$ .

Two simplices  $\sigma$  and  $\tau$  in a complex  $\Sigma$  are called *s-adjacent* if they have a common *s*-face. Hereafter, when we refer to adjacencies into a *d*-complex, we intend to refer to the maximum order adjacencies, i.e. to (d-1)-adjacencies.  $\mathcal{K}_s$   $(s \leq d)$  denotes the set of *s*-faces of  $\Sigma_d$ .

## Simplex orientation

The ordering of 0-skeleton of a simplex implies an orientation of it.

**Definition 3.11 (Ordering of simplex vertices)** The simplex can be oriented according to the even or odd permutation class of its 0-skeleton.  $\Box$ 

The two opposite orientation of a simplex will be denoted as  $+\sigma$  and  $-\sigma$ .

**Definition 3.12 (Coherent orientation)** Two simplices are *coherently oriented* when their common faces have opposite orientation. A complex is *orientable* when all its simplices can be coherently oriented.

It is assumed that:

- 1. the two orientations of a simplex represent its relative interior and exterior;
- 2. the two orientations of an orientable simplicial complex analogously represent the relative interior and exterior of the complex, respectively;
- 3. the boundary of a complex maintains the same orientation of the complex.  $\Box$

The volume associated with an orientation of a simplex (or complex) is positive, while the one associated with the opposite orientation has the same absolute value and opposite sign. It is assumed that the bounded object has positive volume. It is also assumed that either a minus sign or a multiplying factor -1 denote a complementation, i.e. an opposite orientation of the simplex, which can be explicitly obtained by swapping two vertices in its ordered 0-skeleton. For example:

$$+\sigma_3 = \langle \boldsymbol{v}_0, \boldsymbol{v}_1, \boldsymbol{v}_2, \boldsymbol{v}_3 \rangle, \qquad -\sigma_3 = \langle \boldsymbol{v}_1, \boldsymbol{v}_0, \boldsymbol{v}_2, \boldsymbol{v}_3 \rangle.$$

**Definition 3.13 (Volume of a 3-simplex)** The volume of a 3-simplex is a 2-cochain (see Section 3.3) from the cycle of its 2-faces (boundary triangles) to  $\mathbb{R}$ . By definition, a 2-cochain is a map from 2-chains to  $\mathbb{R}$ .

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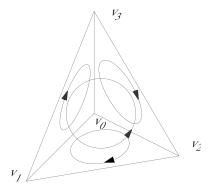


Fig. 3.2 Coherent orientation of the 2-faces of a 3-simplex

#### Facet extraction

The (d-1)-faces of a d-dimensional simplex or complex are also called *facets* since no ambiguity may arise. It is possible to see [?] what follows:

**Theorem 3.1 (Whitney, 1957)** The oriented facets  $\sigma_{d-1}^{(i)}$   $(0 \le i \le d)$  of the oriented d-simplex  $\sigma_d = +\langle \mathbf{v}_0, \mathbf{v}_1, \dots, \mathbf{v}_d \rangle$  are obtained by removing the *i-th* vertex  $\mathbf{v}_i$  from the 0-skeleton of  $\sigma_d$ :

$$\sigma_{d-1}^{(i)} = (-1)^i (\sigma_d - \langle \boldsymbol{v}_i \rangle), \qquad 0 \le i \le d.$$
 (3.3)

The 0-skeleton of  $\sigma_{d-1,(i)}$  is therefore obtained by removing the *i*-th vertex from the 0-skeleton of  $\sigma_d$  and either by swapping a pair of vertices or, better, by inverting the simplex sign, when *i* is odd.

Coding 3.2.5 (Facets of a d-simplex.) A julia function is given here to compute combinatorially all the oriented facets of a simplices arrays of d-simplexes. A precondition is that all Vector{Int64} in the input sequence have the same length.

```
function simplexfacets(simplices)
    @assert hcat(simplices...) isa Matrix
    out = Array{Int64,1}[]
    for simplex in simplices
        for v in simplex
            facet = setdiff(simplex,v)
            push!(out, facet)
    end
end
# remove duplicate facets
return sort(collect(Set(out)))
end
```

If the expr after Qassert is false, the program stops in AssertionError: expr  $\hfill\Box$ 

Coding 3.2.6 (Execution example.) We can start from an array of simplices of the same length and iterate on their facets, to get a whole simplicial complex.

```
FV=[[123],[124],[134],[234]] #=
[[123],[124],[134],[234]] =#
EV = simplexfacets(FV) #=
[[12],[13],[14],[23],[24],[34]] =#
VV = simplexFacets(EV) #=
[[1],[2],[3],[4]] =#
simplex(3,complex=true).C #=
Dict(:c1v => EV, :c3v => [[1,2,3,4], :c0v => VV], :c2v => FV) =#
```

We can also generate directly the whole bounday complex of a multidimensional simplex

Coding 3.2.7 (Generation of boundary complex) We can olso directly generate the Lar boundary complex of a simplex of any degree:

Coding 3.2.8 (Simplex generation in Hpc) Convnersely, the Plasm data structure Hpc contains a single convex cell:

```
SIMPLEX(4) #=
Hpc(MatrixNd(5), Geometry([[0.0, 0.0, 0.0, 0.0], [1.0, 0.0, 0.0, 0.0], [0.0, 1.0, 0.0, 0.0], [0.0, 0.0, 1.0, 0.0], [0.0, 0.0, 1.0, 0.0], [0.0, 0.0, 1.0], hulls=[[1, 2, 3, 4, 5]])) =#
```

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#### Definition 3.14 (Simplicial extrusion)

The prism over a simplex  $\sigma_d = \langle v_0, \dots, v_d \rangle$ , defined as the set  $P_{d+1} := \sigma_d \times [a,b]$ , with  $[a,b] \subset \mathbb{E}$ , will be called  $simplicial\ (d+1)$ -prism. An oriented complex which triangulates  $P_{d+1}$  can be defined combinatorially, by using a closed form formula for its  $\mathcal{K}_{d+1}$  skeleton:

$$\mathcal{K}_{d+1} = \{ \sigma_{d+1,(i)} = (-1)^{id} \langle \boldsymbol{v}_i^a, \boldsymbol{v}_{i+1}^a, \dots, \boldsymbol{v}_d^a, \boldsymbol{v}_0^b, \boldsymbol{v}_1^b, \dots, \boldsymbol{v}_i^b \rangle, \quad 0 \le i \le d \}$$

where  $\mathbf{v}_i^a = (\mathbf{v}_i, a)$  and  $\mathbf{v}_i^b = (\mathbf{v}_i, b)$ .

Closed formulas to triangulate the (d+1)-prism over a d-complex in a time linear with the size of the output, while computing also the d-adjacencies between the resulting (d+1)-simplices, can be found in [?,?].

#### Simplicial grids

In layout design, a grid system provides a framework of intersecting vertical and horizontal lines. Designers use this framework to place and align text, images, and other elements.

Analogously, a *simplicial grid*, is a geometric grid structure made by simplicial cells of same dimension aligned on a layout grid 1D, 2D, 3D, etc. They are mainly used for domain decomposition, where to map coordinate functions in order to create curved structures, i.e., proper curves, curved surfaces, and curved solids.

The whole curved complex is easily generated by mapping the vertices of a Lar grid.

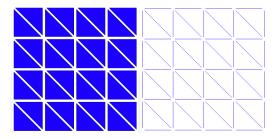


Fig. 3.3 The simplicial 2- and 1-complex (shown here slightly exploded) are generated by extrudecomplex() using the same pattern, both starting from model1D. The 1-complex is derived by the 2-complex using simplexfacets().

#### Coding 3.2.9 (1D Extrusion)

```
V1 = [ 0.0 1.0 2.0 3.0 4.0 5.0 ]
EV = [[1, 2], [2, 3], [3, 4], [4, 5]]
```

```
mod1D = Lar(V1, Dict(:c1v => EV))
V2 = [mod1D.V; zeros(size(mod1D.V,2))']
GL.VIEW(GL.GLExplode(V2, map(x->[x],EV),1.1,1.1,1.1,2,1));
```

The mod2D and mod3D grids are generated by the following code.

#### Coding 3.2.10 (2D Extrusion)

```
mod2D = extrudecomplex(mod1D::Lar, [1, 1, 1, 1])
FV = collect(values(mod2D.C))[1]
EV = simplexfacets(FV)
GL.VIEW(GL.GLExplode(mod2D.V, map(x->[x],EV),1.2,1.2,1.2,4,1));
```

#### Coding 3.2.11 (3D Extrusion)

```
mod3D = extrudecomplex(mod2D::Lar, [1,-2,1])
CV = collect(values(mod3D.C))[1]
FV = simplexfacets(CV)
EV = simplexfacets(FV)
GL.VIEW(GL.GLExplode(mod3D.V,map(x->[x],EV),2,2,2,99,1));
```



#### 3.2.2 Cubical complex and grid

We introduce now the definition and use of multidimensional grids of cuboidal, and the more general  $Cartesian\ product$  of cellular complexes. Such operators, depending on the dimension of their input, may generate either full-dimensional (i.e. solid) output complexes, or lower-dimensional complexes of dimension d embedded in Euclidean n-space, with d < n.

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#### Regular grid

Regular cubical grids of appear on graph paper and may be used in finite element analysis, finite volume methods, finite difference methods, and in general for discretization of parameter spaces.

**Definition 3.15 (Regular grid)** A regular grid is a tessellation of n-dimensional Euclidean space by congruent parallelotopes (e.g. bricks). Its opposite is irregular grid. □

E.g., just think to a mesh of 3D cubes in three-dimensional space for the first case, and to the (non-manifold) framework of boundary polygons of such cubic cells for the second case. In particular, both n-dimensional solid grids of (hyper)-cuboidal cells and their d-dimensional skeletons ( $0 \le d \le n$ ), embedded in  $\mathbb{E}^n$ , are generated by assembling the cells produced by a product of n either 0D or 1D cellular complexes, that in such lowest dimensions coincide with simplicial complexes.

#### 3.2.3 Polyhedral complex

#### 3.2.4 Topological product

In this section, we discuss a solid modeling operation highly useful to generate 3D solids from 2D surfaces, as well as 2-complexes from 1-complexes, and to produce their skeletons taking advantage of one or more 0-complexes.

**Definition 3.16 (Topological product)** A product space is the *Cartesian* product of a family of topological spaces equipped with a natural topology.

The natural topology on a subset of a topological space, in our case  $\mathbb{E}^d$ , is the relative topology (or subspace topology). The product of two cell complexes can be made into a cell complex as we show in the following.

**Definition 3.17 (Product of cell complexes)** If X and Y are m- and n-complexes in  $\mathbb{E}^m$  and  $\mathbb{E}^n$ , respectively,  $X \times Y \subset \mathbb{E}^{m+n}$  is a cell complex in which each cell is a product of a cell in X and a cell in Y, endowed with the relative topology in  $\mathbb{E}^{m+n}$ .

#### 3.2.5 Cell complex product

**Definition 3.18 (larprod)** The larprod function takes as input a pair of Lar models and returns the model of their *Cartesian product*. Since the Lar type is a pair (*geometry, topology*), the output of larprod( lar1, lar2 ) is the *topological product* of the input Lar values.

#### 3.3 Chain complex

#### 3.3.1 Linear chain spaces

#### 3.3.2 Linear chain operators

#### Matrix-Vector Products as Linear Combinations

Left-multiply a matrix A by a row vector x, i.e. xA, is a linear combination of A rows with x elements as scalars, and produces a row vector. Analogously, right-multiply a matrix A by a column vector x, i.e. Ax, is a linear combination of A columns, and produces a column vector.

In this book we often use a pair of characters from  $\{V, E, F\}$ , for instance EF, to denote the mapping  $F \to E$  from 2-chains in F to 1-chains in E (see Section 3.1).

Coding 3.3.1 (Construction of operator matrix) [ $\partial_2$ ]. With some abuse of language (removed after Section ??), we build the sparse matrix of EF  $\equiv \partial_2$ : F  $\rightarrow$  E.

It may be interesting to make some notation about this matrix. Remember that, by right multiplication, any mapping matrix sends the space of columns (2-chains: triangles here) to the space of rows (1-chains: edges here). With any simplicial complex we have two 1s on each row and three 1s on each column. They characterize the two faces incident on the edge, and the three edges incident on the face, respectively.

Coding 3.3.2 (How to use the linear operators.) A boundary operator, which is a linear map, sends every vector in domain space (the span of the matrix columns) to its image vector in target space (the span of the matrix rows).

In particular, the matrix of a linear operator—coordinate representation of it with respect to the chosen bases—send the coordinate representation of a domain vector to its image in target space. Both are binary vectors, as will be clear after Section ??. E.g., face[4] has boundary edges 3, 4, and 12 (look for 1s positions in row 5 above).

#### 3.4 Cochain integration (surface, volume, inertia)

Coding 3.4.1 (Volume of a cube from boundary triangles) . The variable obj first contains the memory pointer to an object of type Hpc, then to an object of type Lar. Here, obj.C gives the dictionary C of its cell complex, where C[:FV] denotes the array of boundary faces (triangles).

```
julia> obj = SIMPLEX(3)
Hpc(MatrixNd(4), Geometry([[0.0, 0.0, 0.0], [1.0, 0.0, 0.0],
     [0.0, 1.0, 0.0], [0.0, 0.0, 1.0]], hulls=[[1, 2, 3, 4]]))
julia> obj = LAR(SIMPLEX(3))
Lar(3, 3, 4, [0.0 1.0 0.0 0.0; 1.0 0.0 0.0 0.0; 0.0 0.0 0.0
    1.0], Dict{Symbol, AbstractArray}(:CV => [[1, 2, 3, 4]], :FV => [[1, 2, 3], [2, 3, 4], [1, 3, 4], [1, 2, 4]], :EV => [[2, 3], [1, 3], [1, 2], [3, 4], [2, 4], [1, 4]]))
FV = obj.C[:FV] #=
4-element Vector{Vector{Int64}}:
 [1, 2, 3]
 [1, 2, 4]
 [1, 3, 4]
 [2, 3, 4]
               =#
P = (obj.V, FV); VOLUME(P)
0.16666666666666666
volume(P) * 6
1.0
```

## Part II Dimension-independent Modeling

# Chapter 4 Geometric models

Julia Plasm is the best choice to develop geometric models for Building Information Modeling (BIM) and Computer-Aided Design (CAD). We ported the functional language to Julia for better supporting design, model generation, and visualization of geometric objects. Geometric models specify the physical appearance of Architecture, Engineering, and Construction products at any scale, from structural and envelope components to whole buildings and built environments, and are used for design, tender, contract, and collaboration. In this chapter we introduce the great expressive power of Plasm parametric functions and the simple methods of parametric assemblies, where objects itself can be used as actual parameters. We show also that Plasm offers a very general mechanism (Julia dictionaries) to characterize and export models with colors, textures, materials, and so on. Plasm can be even embedded in the Jupiter platform to document the design choices step-by-step.

- 4.1 Introduction to geometric objects
- 4.2 Plasm parametric primitives
- 4.3 Hierarchical assembly of geometric objects
- 4.4 Attach properties to geometry
- 4.5 Design documentation (Jupyter notebooks)
- 4.6 Export geometry

# Chapter 5 Symbolic modeling with Julia Plasm

Symbolic modeling is a semantic approach to knowledge representation and processing. The symbolic approach to knowledge representation and processing uses names to define the meaning of represented knowledge explicitly. The geometric knowledge is described here by Julia's names, which are given to functionals, functions, formal and actual parameters, and finally to objects, fields, classes, attributes, methods, relations, etc. In this chapter, we give many examples of high-level Plasm programming, from topological, linear, and affine operators, to topological mapping of complexes and grids to generate linearized approximation of curved manifold of intrinsic dimensions 1, 2, and 3. i.e., depending on such number of parameters; say, curves, surfaces, thin, and bulk solids.

- 5.1 Primitive generators
- 5.2 Plasm topological operators
- 5.3 Linear and affine operators
- 5.4 Manifold mapping
- 5.5 Predefined Plasm functions
- 5.6 Curve, surface, and solid methods

## Chapter 6 Product assembly structure

Hierarchical models of assemblies are generated by aggregating subassemblies, each defined in its local coordinate system and relocated by affine transformations. This operation may be repeated hierarchically, with some subassemblies defined by aggregating simpler parts, and so on, until we get a set of elementary components that cannot be further disaggregated. Two main advantages can be found in hierarchical modeling. Each elementary part and each assembly, at every hierarchical level, are defined independently from each other, using local coordinate frames suitably chosen to make their definition easier. Furthermore, only one copy of each component is used, and it can be instanced in different locations and orientations, depending on how many times it is needed. In this chapter, after an overview of solid modeling data structures, the models of the Plasm dataset are discussed.

- 6.1 Hierarchical ssembly definition
- 6.2 Data structures in solid modeling
- 6.3 Structure in PHIGS and Plasm
- 6.4 Julia Plasm data structures
- 6.4.1 Hierarchical Polyhedral Complex (HPC)
- 6.4.2 Linear Algebraic Representation (LAR)
- 6.4.3 Geometric DataSet (GEO)

# Chapter 7 Space arrangements

The chapter introduces computational topology algorithms to discover the two-dimensional (2D)/3D space partition induced by a collection of geometric objects of dimension (1D)/2D, respectively. The data structures needed for such computational program are sparse arrays and their standard algebraic operations. In this chapter, we introduce a novel approach to solid modeling based on piecewise-linear algebraic topology, that allows to treat rather general cellular complexes, with cells homeomorphic to polyhedra, i.e., to triangulable spaces, and hence possibly non-convex and multiply connected. The notions we deal with include geometric complexes, linear spaces of chains and cochains, the chain complex of linear operators between pairs of spaces, and their compositions. The discussion is restricted to piecewise-linear topology and to space dimensions less or equal to three.

- 7.1 Space partition and enumeration
- 7.2 Cellular and boundary models
- 7.3 Arrangements and Lattices
- 7.4 2D and 3D Examples

## Chapter 8 Boolean solid algebras

Engineers conceived Solid Modeling as a rigorous and universal language for geometry-based engineering. Mathematically, all solid models are computer representations of elements of some algebraic systems that are constructed and maintained by algorithms corresponding to the operations in such algebra. This chapter, in particular, focuses on the algebraic relationships between CSG (Constructive Solid Geometry) and boundary representations of PL polyhedra. We show that the (regularized) arrangement of a given set of spatially instanced primitive shapes is isomorphic to the finite Boolean algebra of regularized sets containing all possible CSG representations with the same primitives. In particular, we show that the boundary evaluation of any CSG representation with a finite number of primitives reduces algebraically to a composition of Boolean operations on bit strings, sparse matrix multiplication, and point membership tests, and show significant examples.

- 8.1 Constructive Solid Geometry (CSG)
- 8.2 Atoms and Generators
- 8.3 Finite Boolean Algebras
- 8.4 Computational Pipeline

# $\begin{array}{c} {\rm Part~III} \\ {\rm Polyhedral~Modeling~in~AEC} \end{array}$

# Chapter 9 Building Information Modeling (BIM)

The construction industry is nowadays undergoing its very own digital revolution, helped by its unique way of working with BIM systems whose elements are the digital prototypes of physical elements like columns, windows, walls, doors, and stairs. In this chapter, we outline the conceptual foundations and the down of the BIM attitude of the construction industry, discuss the birth of concepts like building objects and the standardized taxonomy of building systems as set of subsystems devoted to the fulfillment of physical requirements of space with suitable performance levels. In particular, we show by example how Julia Plasm, relying on its understand of topology, may automatically transform the preliminary shape design of spaces into the generic geometry of building subsystems (skeleton, envelope, internal partitions, vertical communications) and elements.

- 9.1 BIM history (Chuck Eastman, ...)
- 9.2 Building taxonomy (UNI 9838)
- 9.3 Building envelope
- 9.4 Building skeleton
- 9.5 Construction Process Modeling

### Chapter 10 Industry Foundation Classes (IFC)

The Industry Foundation Classes (IFC) are an open international standard for sharing Building Information Model (BIM) data. IFC is a non-proprietary file format used to exchange building information models. IFC files can share data between different software applications, making it possible for multiple stakeholders to collaborate on BIM projects. Of course, we are mostly interested in the IFC transport of shape information. In this chapter, we, therefore, investigate the Julia Plasm ability to implement in a few rows of parametric code the classes and attributes of the IfcShapeRepresentation entity for the various types of shape representation, including PointCloud, Curve, Surface, GeometricSet, GeometricCurveSet, Annotation2D, SurfaceModel, Tessellation, SolidModel, SweptSolid, AdvancedSweptSolid, Brep, CSG, Clipping, and BoundingBox.

- 10.1 Simple introduction to IFC
- 10.2 Data scheme for BIM collaboration
- 10.3 IfcShapeRepresentation (IFC 4.3.x: 8.18.3.14)
- 10.3.1 Representation identifiers
- 10.3.2 Representation types
- 10.3.3 Representation Examples
- 10.4 Plasm parametric programming to IFC

## Part IV Geometry from Point Cloud